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7931	79877194 (6551, 6552)	Novel Protein sim. GBank gj 3294482 (AF001386) - unknown [Salmonella typhimurium]		UNCLASSIFIED	1023
7932	94747030 (18049, 18050)	Novel Protein sim. GBank gj 3299896 (AF020390) - beta-galactosidase [Lycopersicon esculentum]		UNCLASSIFIED	1003
7933	20608435 (4051, 4052)	Novel Protein sim. GBank gj 3319723[emb CAA19889] - (AL031035) putative integral membrane protein [Streptomyces coelicolor]		UNCLASSIFIED	1004
7934	87457287 (18551, 18552)	Novel Protein sim. GBank gj 3319730[emb CAA19896] - (AL031035) transcriptional regulatory protein [Streptomyces coelicolor]		UNCLASSIFIED	1025
7935	55652892 (14969, 14970)	Novel Protein sim. GBank gj 3327044[db BAA31590] - (AB014515) KIAA0615 protein [Homo sapiens]		UNCLASSIFIED	1000
7936	17253192 (10421, 10422)	Novel Protein sim. GBank gj 3327088[db BAA31612] - (AB014537) KIAA0637 protein [Homo sapiens]		UNCLASSIFIED	1023
7937	20710076 (7819, 7820)	Novel Protein sim. GBank gj 3328936 (AE001323) - Lipocate Protein Ligase [Chlamydia trachomatis]		UNCLASSIFIED	1022
7938	78287717 (13641, 13642)	Novel Protein sim. GBank gj 3329465 (AF064553) - NSD1 protein [Mus musculus]		UNCLASSIFIED	1050
7939	80078084 (15935, 15936)	Novel Protein sim. GBank gj 3334134[sp Q12495 CAC1_YEAST - CHROMATIN ASSEMBLY FACTOR 1 P90 SUBUNIT (CAF-1 90 KD SUBUNIT) (RAP1 LOCALIZATION FACTOR 2)]		UNCLASSIFIED	1008, 1010, 1038, 1039
7940	87138194 (15973, 15974)	Novel Protein sim. GBank gj 3334526[emb CAA16138] - (AL021306) predicted using FGENEH [Homo sapiens]		UNCLASSIFIED	1007
7941	79828340 (15471, 15472)	Novel Protein sim. GBank gj 3335359 (AC003028) - unknown protein [Arabidopsis thaliana]		UNCLASSIFIED	1003, 1008, 1017, 1023, 1026

7942	12892835 (14229, 14230)	Novel Protein sim. GBank gi 3342232 (U93160) - major membrane glycoprotein [Cercopithecine herpesvirus 15]		UNCLASSIFIED	1031
7943	25256839 (4563, 4564)	Novel Protein sim. GBank gi 3342713 (U22894) - alpha glucosidase [Streptomyces lividans]		UNCLASSIFIED	1008
7944	87451658 (4057, 4058)	Novel Protein sim. GBank gi 3367534 (AC004392) - Strong similarity to coarmer alpha subunit (HEPCOP) homolog gb U24105 from Homo sapiens. [Arabidopsis thaliana]		UNCLASSIFIED	1025
7945	30238805 (4685, 4686)	Novel Protein sim. GBank gi 3378278 (AF079317) - putative aromatic efflux pump membrane protein [Sphingomonas aromaticivorans]		UNCLASSIFIED	1008
7946	10063314 (4141, 4142)	Novel Protein sim. GBank gi 3411182 (AF076240) - putative regulatory protein MocR [Rhizobium leguminosarum bv. viciae]		UNCLASSIFIED	1017
7947	20728168 (17177, 17178)	Novel Protein sim. GBank gi 3414725 (AF047693) - multidrug resistance efflux pump homolog PmrA [Pseudomonas aeruginosa]		UNCLASSIFIED	1006
7948	80054207 (13109, 13110)	Novel Protein sim. GBank gi 3417424[emb CAA20312] - (AL031261) putative transport protein [Schizosaccharomyces pombe]		UNCLASSIFIED	1004
7949	94631686 (21753, 21754)	Novel Protein sim. GBank gi 3449276[emb CAA20420] - (AL031317) putative dehydrogenase [Streptomyces coelicolor]		UNCLASSIFIED	1006, 1012, 1031, 1034
7950	79579867 (17353, 17354)	Novel Protein sim. GBank gi 3451460[emb CAA20477.1] - (AL031349) zinc-finger protein [Schizosaccharomyces pombe]		UNCLASSIFIED	1004
7951	78768130 (6205, 6206)	Novel Protein sim. GBank gi 3452497[emb CAA76854] - (Y17796) ketol-acid reductoisomerase [Pisum sativum]		UNCLASSIFIED	1003, 1017, 1044
7952	57293806 (9705, 9706)	Novel Protein sim. GBank gi 348457[pir A44154] - iron-responsive element-binding protein - rat		UNCLASSIFIED	1008, 1016, 1022, 1038, 1044

7953	85527214 (20607, 20608)	Novel Protein sim. GBank gj 3522880 (U32222) - P2 J homolog; baseplate or base of tail fibre [Bacteriophage 186]		UNCLASSIFIED	1038
7954	50875118 (14859, 14860)	Novel Protein sim. GBank gj 3550295 emb CAA08912 - (AJ009947) putative ATPase [Homo sapiens]		UNCLASSIFIED	1036
7955	80060356 (12057, 12058)	Novel Protein sim. GBank gj 3581849 emb CAA20805 - (AL031541) putative phenylalanyl-tRNA synthetase beta chain [Streptomyces coelicolor]		UNCLASSIFIED	1024
7956	21432731 (2775, 2776)	Novel Protein sim. GBank gj 3622 emb CAA32032 - (X13793) CYP1 protein (AA 1-1483) [Saccharomyces cerevisiae]		UNCLASSIFIED	1010
7957	32153290 (10283, 10284)	Novel Protein sim. GBank gj 3641340 gb AAC36351 - (AF090329) cyclohexanone monooxygenase homolog [Pseudomonas fluorescens]		UNCLASSIFIED	1039
7958	79470508 (11609, 11610)	Novel Protein sim. GBank gj 3641340 gb AAC36351 - (AF090329) cyclohexanone monooxygenase homolog [Pseudomonas fluorescens]		UNCLASSIFIED	1022
7959	30643710 (8383, 8384)	Novel Protein sim. GBank gj 3650033 (AC005396) - unknown protein [Arabidopsis thaliana]		UNCLASSIFIED	1034
7960	11359161 (17651, 17652)	Novel Protein sim. GBank gj 3650037 (AC005396) - auxin-responsive GH3-like protein [Arabidopsis thaliana]		UNCLASSIFIED	1058
7961	29459767 (12617, 12618)	Novel Protein sim. GBank gj 3659692 (AF068748) - sphingosine kinase [Mus musculus]		UNCLASSIFIED	1026
7962	27786197 (11737, 11738)	Novel Protein sim. GBank gj 3651614 (AF093142) - aconitase [Aspergillus terreus]		UNCLASSIFIED	1022
7963	78077206 (21067, 21068)	Novel Protein sim. GBank gj 3668151 emb CAA21108.1 - (AL031764) putative FAD synthetase [Schizosaccharomyces pombe]		UNCLASSIFIED	1029

7964	80027018 (4693, 4694)	Novel Protein sim. GBank gil3676737 gb AAC64273.1 AAC6427 - (AF086832) unknown [Streptomyces coelicolor]		UNCLASSIFIED	1004, 1006, 1010, 1022, 1034
7965	20732951 (6773, 6774)	Novel Protein sim. GBank gil3676737 gb AAC64273.1 AAC6427 - (AF086832) unknown [Streptomyces coelicolor]		UNCLASSIFIED	1006
7966	78753197 (9569, 9570)	Novel Protein sim. GBank gil3695123 (AF081148) - CL2AA [Rattus norvegicus]		UNCLASSIFIED	1000, 1022
7967	20294748 (6849, 6850)	Novel Protein sim. GBank gil3724125 emb CAA11905] - (AJ224340) maltosephosphorylase [Lactobacillus sanfrancisco]		UNCLASSIFIED	1034
7968	30390483 (6647, 6648)	Novel Protein sim. GBank gil3724134 emb CAA09789.1] - (AJ011810) HASP B1 protein [Leishmania donovani]		UNCLASSIFIED	1008, 1034
7969	80052628 (2493, 2494)	Novel Protein sim. GBank gil3738200 emb CAA21292] - (AL031855) putative vacuolar membrane protein [Schizosaccharomyces pombe]		UNCLASSIFIED	1010, 1024
7970	86678720 (801, 802)	Novel Protein sim. GBank gil3738309 (AC005309) - unknown protein [Arabidopsis thaliana]		UNCLASSIFIED	1007
7971	82114121 (10699, 10700)	Novel Protein sim. GBank gil3747107 (AF095741) - unknown [Rattus norvegicus]		UNCLASSIFIED	1016, 1022, 1026
7972	78527551 (10701, 10702)	Novel Protein sim. GBank gil3747111 (AF095641) - MTN3 homolog [Arabidopsis thaliana]		UNCLASSIFIED	1003
7973	11309497 (6377, 6378)	Novel Protein sim. GBank gil3758880 emb CAA09871] - (AJ011927) fructose- 1,6-bisphosphate aldolase [Pseudomonas stutzeri]		UNCLASSIFIED	1010
7974	25259590 (14051, 14052)	Novel Protein sim. GBank gil3763 emb CAA34707] - (X16727) HAP4 (AA 1- 554) [Saccharomyces cerevisiae]		UNCLASSIFIED	1026

7975	11076716 (11969, 11970)	Novel Protein sim. GBank gij3789932 (AF090142) - lipase precursor [Staphylococcus epidermidis]		UNCLASSIFIED	1010
7976	78902638 (14901, 14902)	Novel Protein sim. GBank gij3800951 gbjAAC68959.1 - (AF100657) No definition line found [Caenorhabditis elegans]		UNCLASSIFIED	1022, 1044
7977	14218581 (11153, 11154)	Novel Protein sim. GBank gij3805990 (AF025969) - unknown [Helicobacter pylori]		UNCLASSIFIED	1053
7978	10840441 (10611, 10612)	Novel Protein sim. GBank gij3845248 (AE001410) - hypothetical protein [Plasmodium falciparum]		UNCLASSIFIED	1017
7979	79602734 (12381, 12382)	Novel Protein sim. GBank gij3850063 embjCAA11566 - (AJ223830) ARE1 [Rattus norvegicus]		UNCLASSIFIED	1034
7980	20466372 (12633, 12634)	Novel Protein sim. GBank gij3850752 embjCAA05698 - (AJ002735) initiation factor IF2-alpha [Klebsiella oxytoca]		UNCLASSIFIED	1010
7981	57302724 (4675, 4676)	Novel Protein sim. GBank gij3859570 gbjAAC72848.1 - (AF098753) unknown [Oryza sativa]		UNCLASSIFIED	1010, 1053
7982	20729698 (1487, 1488)	Novel Protein sim. GBank gij3860997 embjCAA14897 - (AJ235271) unknown [Rickettsia prowazekii]		UNCLASSIFIED	1006
7983	20474205 (3719, 3720)	Novel Protein sim. GBank gij3861224 embjCAA15124 - (AJ235272) VALYL- TRNA SYNTHETASE (valS) [Rickettsia prowazekii]		UNCLASSIFIED	1010
7984	11101579 (5259, 5260)	Novel Protein sim. GBank gij3868935 dbjBAA34293 - (AB015023) MurC [Corynebacterium glutamicum]		UNCLASSIFIED	1006
7985	80231085 (4885, 4886)	Novel Protein sim. GBank gij3873958 embjCAB03821 - (Z81458) similar to transposable element [Caenorhabditis elegans]		UNCLASSIFIED	1058

7986	10857565 (22103, 22104)	Novel Protein sim. GBank gi 3876233 emb CAA90360 - (Z50044) similar to phenylalanyl-tRNA synthetase; cDNA EST EMBL:T01401 comes from this gene; cDNA EST yk303c5.3 comes from this gene; cDNA EST yk303c5.5 comes from this gene; cDNA EST yk452d5.3 comes from this gene; cDNA EST yk...		UNCLASSIFIED	1006
7987	30659749 (19625, 19626)	Novel Protein sim. GBank gi 3877579 emb CAB05214.1 - (Z82271) Similarity to Mouse kinensin-like protein KIF4 (SW:P33174); cDNA EST EMBL:D27320 comes from this gene; cDNA EST EMBL:D27322 comes from this gene; cDNA EST EMBL:D27321 comes from this gene; cDNA EST EMBL:D35764 comes ...		UNCLASSIFIED	1022
7988	11310115 (3933, 3934)	Novel Protein sim. GBank gi 3877645 emb CAB05741 - (Z83230) cDNA EST yk355g3.5 comes from this gene [Caenorhabditis elegans]		UNCLASSIFIED	1022, 1039
7989	90995295 (14129, 14130)	Novel Protein sim. GBank gi 3877645 emb CAB05741 - (Z83230) cDNA EST yk355g3.5 comes from this gene [Caenorhabditis elegans]		UNCLASSIFIED	1020, 1044
7990	28987108 (5845, 5846)	Novel Protein sim. GBank gi 3880413 emb CAB03470 - (Z81137) Similarity to Yeast YIP1 protein (SW:P53039); cDNA EST EMBL:T01608 comes from this gene; cDNA EST EMBL:C07393 comes from this gene; cDNA EST EMBL:C07550 comes from this gene; cDNA EST EMBL:C08746 comes from this gene; ...		UNCLASSIFIED	1023

7991	20628075 (13809, 13810)	Novel Protein sim. GBank gi 3880428 emb CAA91394 - (Z66521) similar to pre-mRNA splicing factor like protein; cDNA EST EMBL:T01363 comes from this gene; cDNA EST EMBL:D65797 comes from this gene; cDNA EST EMBL:D72951 comes from this gene; cDNA EST EMBL:D69365 comes from this ge...		UNCLASSIFIED	1010
7992	86692011 (10901, 10902)	Novel Protein sim. GBank gi 3882321 dbj BAA34520.1 - (AB018343) KIAA0800 protein [Homo sapiens]		UNCLASSIFIED	1025
7993	79397657 (9309, 9310)	Novel Protein sim. GBank gi 3882325 dbj BAA34522.1 - (AB018345) KIAA0802 protein [Homo sapiens]		UNCLASSIFIED	1016, 1034
7994	94234704 (19263, 19264)	Novel Protein sim. GBank gi 3913029 sp P94967 ALR_MYCSM - ALANINE RACEMASE		UNCLASSIFIED	1003, 1010
7995	20711506 (11735, 11736)	Novel Protein sim. GBank gi 3913102 sp O50175 ASTB_PSEAE - SUCCINYLARGININE DIHYDROLASE		UNCLASSIFIED	1006
7996	79836788 (14783, 14784)	Novel Protein sim. GBank gi 3913102 sp O50175 ASTB_PSEAE - SUCCINYLARGININE DIHYDROLASE		UNCLASSIFIED	1008
7997	25136730 (12599, 12600)	Novel Protein sim. GBank gi 3913656 sp O52256 FLHF_PSEPU - FLAGELLAR BIOSYNTHESIS PROTEIN FLHF (FLAGELLA ASSOCIATED GTP-BINDING PROTEIN)		UNCLASSIFIED	1008
7998	20288848 (10675, 10676)	Novel Protein sim. GBank gi 3914317 sp Q19189 PEXC_CAEEL - PUTATIVE PEROXIN-12		UNCLASSIFIED	1034
7999	78729545 (19001, 19002)	Novel Protein sim. GBank gi 3914643 sp O33016 RIMM_MYCLE - PROBABLE 16S RNA PROCESSING PROTEIN RIMM		UNCLASSIFIED	1008

8000	16419204 (3859, 8860)	Novel Protein sim. GBank gil3915530 sp Q12496 YOJ8_YEAST - HYPOTHETICAL 118.4 KD PROTEIN IN WRS1- PKH2 INTERGENIC REGION	UNCLASSIFIED	1003
8001	306339016 (19589, 19590)	Novel Protein sim. GBank gil3915530 sp Q12496 YOJ8_YEAST - HYPOTHETICAL 118.4 KD PROTEIN IN WRS1- PKH2 INTERGENIC REGION	UNCLASSIFIED	1022
8002	78770735 (2001, 2002)	Novel Protein sim. GBank gil3915566 sp O32156 YUO_BACSU - HYPOTHETICAL ABC TRANSPORTER EXTRACELLULAR BINDING PROTEIN YUO PRECURSOR	UNCLASSIFIED	1017
8003	28991603 (5833, 5834)	Novel Protein sim. GBank gil3915759 sp P36044 MNNA4_YEAST - MNNA4 PROTEIN	UNCLASSIFIED	1044
8004	78461435 (16941, 16942)	Novel Protein sim. GBank gil3915986 sp P37908 YFJD_ECOLI - HYPOTHETICAL 46.4 KD PROTEIN IN FFH- GRPE INTERGENIC REGION	UNCLASSIFIED	1026
8005	11694498 (16815, 16816)	Novel Protein sim. GBank gil3915990 sp P54585 YHCA_BACSU - HYPOTHETICAL 58.3 KD PROTEIN IN GLPD- CSPB INTERGENIC REGION	UNCLASSIFIED	1034
8006	80028400 (2489, 2490)	Novel Protein sim. GBank gil3915993 sp P76250 YEAT_ECOLI - HYPOTHETICAL TRANSCRIPTIONAL REGULATOR IN GAPA-RND INTERGENIC REGION	UNCLASSIFIED	1008, 1022
8007	19637574 (4223, 4224)	Novel Protein sim. GBank gil3924708 emb CAA84646 - (Z35597) Weak similarity with sea squirt nidogen precursor protein (blastp score 71); cDNA EST EMBL:T02069 comes from this gene; cDNA EST EMBL:D76135 comes from this gene; cDNA EST EMBL:D73147 comes from this gene; cDNA EST EMB...	UNCLASSIFIED	1054

8008	17664545 (4937, 4938)	Novel Protein sim. GBank gil3924708 embjCAA84646 - (Z35597) Weak similarity with sea squirt nidogen precursor protein (blastp score 71); cDNA EST EMBL:T02069 comes from this gene; cDNA EST EMBL:D76135 comes from this gene; cDNA EST EMBL:D73147 comes from this gene; cDNA EST EMB...		UNCLASSIFIED	1008
8009	78385723 (11405, 11406)	Novel Protein sim. GBank gil3928766 dbjBAA34704 - (AB020491) cathepsin L-like tick cysteine proteinase B [Haemaphysalis longicornis] Novel Protein sim. GBank gil3935187 (AC004557) - F17L21.30 [Arabidopsis thaliana]		UNCLASSIFIED	1026
8010	13502044 (14209, 14210)	Novel Protein sim. GBank gil3928766 dbjBAA34704 - (AB020491) cathepsin L-like tick cysteine proteinase B [Haemaphysalis longicornis]		UNCLASSIFIED	1027
8011	23308587 (22681, 22682)	Novel Protein sim. GBank gil3935187 (AC004557) - F17L21.30 [Arabidopsis thaliana]		UNCLASSIFIED	1053
8012	25322989 (20611, 20612)	Novel Protein sim. GBank gil3947608 embjCAA19460.1 - (AL023828) Y17G7B.13 [Caenorhabditis elegans]		UNCLASSIFIED	1003
8013	17300330 (13315, 13316)	Novel Protein sim. GBank gil396348 (U00006) - homoserine transsuccinylase [Escherichia coli]		UNCLASSIFIED	1044
8014	70336470 (9985, 9986)	Novel Protein sim. GBank gil3980411 (AC004561) - putative proline-rich protein [Arabidopsis thaliana]		UNCLASSIFIED	1023, 1049
8015	87122170 (3345, 3346)	Novel Protein sim. GBank gil3980412 (AC004561) - pumilio-like protein [Arabidopsis thaliana]		UNCLASSIFIED	1025
8016	20466868 (7279, 7280)	Novel Protein sim. GBank gil4001725 dbjBAA35095 - (AB015981) MnhA [Staphylococcus aureus]		UNCLASSIFIED	1010
8017	39517664 (22689, 22690)	Novel Protein sim. GBank gil400324 sp P31179 NADA_CYAPA - QUINOLINATE SYNTHETASE A		UNCLASSIFIED	1006
8018	6325410 (4993, 4994)	Novel Protein sim. GBank gil4006920 embjCAB16815.1 - (Z99708) actin interacting protein [Arabidopsis thaliana]		UNCLASSIFIED	1058

8019	11083825 (16171, 16172)	Novel Protein sim. GBank gil4007680[emb]CAA22366] - (AL034443) putative oxidoreductase [Streptomyces coelicolor]		UNCLASSIFIED	1024
8020	27962805 (13965, 13966)	Novel Protein sim. GBank gil401086[sp]Q02793[SK18_YEAST - ANTIVIRAL PROTEIN SK18]		UNCLASSIFIED	1034
8021	78729716 (16117, 16118)	Novel Protein sim. GBank gil401087[sp]Q02100[SK01_YEAST - CRE- BINDING BZIP PROTEIN SK01]		UNCLASSIFIED	1022, 1026
8022	80050523 (10589, 10590)	Novel Protein sim. GBank gil401223[sp]P06969[TTK_ECOLI - TTK PROTEIN]		UNCLASSIFIED	1004
8023	78776707 (20435, 20436)	Novel Protein sim. GBank gil401441[sp]P31382[PMT2_YEAST - DOLICHYL- PHOSPHATE-MANNOSE--PROTEIN MANNOSYLTRANSFERASE 2]		UNCLASSIFIED	1009, 1017
8024	17958538 (273, 274)	Novel Protein sim. GBank gil401561[sp]P31120[MRSA_ECOLI - MRSA PROTEIN]		UNCLASSIFIED	1016
8025	11607959 (20371, 20372)	Novel Protein sim. GBank gil401562[sp]P27432[YICE_ECOLI - HYPOTHETICAL 48.9 KD PROTEIN IN GLTS- SELC INTERGENIC REGION]		UNCLASSIFIED	1004
8026	78751746 (10815, 10816)	Novel Protein sim. GBank gil4050087 (AF109907) - S164 [Homo sapiens]		UNCLASSIFIED	1026
8027	16303494 (15571, 15572)	Novel Protein sim. GBank gil4050087 (AF109907) - S164 [Homo sapiens]		UNCLASSIFIED	1022
8028	78478238 (11727, 11728)	Novel Protein sim. GBank gil4062975[dbj]BAA36206.1] - (AB017138) delta subunit of malonate decarboxylase [Pseudomonas putida]		UNCLASSIFIED	1003, 1006
8029	79412280 (14065, 14066)	Novel Protein sim. GBank gil4063042 (AF068065) - GP900; mucin-like glycoprotein [Cryptosporidium parvum]		UNCLASSIFIED	1001, 1022, 1026

8030	85531233 (16933, 16934)	Novel Protein sim. GBank gi 4063795 dbj BAA36249.1 - (AB008550) orf22; similar to F1 gene of P2 [Pseudomonas aeruginosa phage phi CTX]		UNCLASSIFIED	1038
8031	30277206 (3741, 3742)	Novel Protein sim. GBank gi 4088 emb CAA49508 - (X69881) ORF2 [Saccharomyces cerevisiae]		UNCLASSIFIED	1026
8032	79598341 (10427, 10428)	Novel Protein sim. GBank gi 4097459 (U62039) - reverse transcriptase [Elephantulus edwardii]		UNCLASSIFIED	1004
8033	11759315 (11543, 11544)	Novel Protein sim. GBank gi 4097495 (U62910) - kexin [Pneumocystis carinii]		UNCLASSIFIED	1024
8034	80195326 (2569, 2570)	Novel Protein sim. GBank gi 4101574 (AF004876) - 54TmP [Homo sapiens]		UNCLASSIFIED	1003, 1006, 1022, 1034
8035	79571122 (7587, 7588)	Novel Protein sim. GBank gi 4102881 (AF017250) - vitellogenin precursor [Oreochromis aureus]		UNCLASSIFIED	1001, 1026, 1032
8036	79911377 (11099, 11100)	Novel Protein sim. GBank gi 4105359 (AF045240) - unknown [Staphylococcus epidermidis]		UNCLASSIFIED	1017
8037	86380904 (10031, 10032)	Novel Protein sim. GBank gi 4106673 emb CAA22613 - (AL035064) queuine tRNA-ribosyltransferase [Schizosaccharomyces pombe]		UNCLASSIFIED	1053
8038	5496348 (939, 940)	Novel Protein sim. GBank gi 4115936 gb AAD03446.1 - (AF118223) No definition line found [Arabidopsis thaliana]		UNCLASSIFIED	1058
8039	17698610 (12979, 12980)	Novel Protein sim. GBank gi 414081 emb CAA53202 - (X75449) RAS GTPase-activating protein [Saccharomyces cerevisiae]		UNCLASSIFIED	1044
8040	80218678 (22771, 22772)	Novel Protein sim. GBank gi 414081 emb CAA53202 - (X75449) RAS GTPase-activating protein [Saccharomyces cerevisiae]		UNCLASSIFIED	1006, 1024, 1025, 1026, 1034, 1038, 1039

8041	80499668 (11509, 11510)	Novel Protein sim. GBank gl 4154082 emb CAA22735 - (AL035161) putative aminoacylase [Streptomyces coelicolor]		UNCLASSIFIED	1012, 1024, 1031, 1034
8042	14206648 (13361, 13362)	Novel Protein sim. GBank gl 4154536 (AE001442) - putative [Helicobacter pylori J99]		UNCLASSIFIED	1034
8043	13883873 (20997, 20998)	Novel Protein sim. GBank gl 4154819 (AE001466) - putative [Helicobacter pylori J99]		UNCLASSIFIED	1053
8044	14401769 (13323, 13324)	Novel Protein sim. GBank gl 4155124 (AE001490) - putative [Helicobacter pylori J99]		UNCLASSIFIED	1053
8045	13883854 (17751, 17752)	Novel Protein sim. GBank gl 4155692 (AE001537) - putative Outer membrane protein [Helicobacter pylori J99]		UNCLASSIFIED	1053
8046	20422258 (19605, 19606)	Novel Protein sim. GBank gl 4155750 (AE001542) - D-LACTATE DEHYDROGENASE [Helicobacter pylori J99]		UNCLASSIFIED	1024
8047	37800193 (13271, 13272)	Novel Protein sim. GBank gl 4160314 emb CAA22785 - (AL035212) putative AraC family transcriptional regulator [Streptomyces coelicolor]		UNCLASSIFIED	1012
8048	79574464 (6625, 6626)	Novel Protein sim. GBank gl 4160344 emb CAA22812 - (AL035216) probable involvement in ergosterol biosynthesis [Schizosaccharomyces pombe]		UNCLASSIFIED	1031
8049	79822297 (10383, 10384)	Novel Protein sim. GBank gl 4160660 gb AAD05269.1 - (AF109386) protocatechuate 3,4-deoxygenase beta subunit; PcaH; 3,4-PCD [Streptomyces sp. 2065]		UNCLASSIFIED	1034
8050	28776278 (19365, 19366)	Novel Protein sim. GBank gl 416288 dbj BAA03957 - (D16506) ORF1 [Saccharomyces cerevisiae]		UNCLASSIFIED	1006, 1008, 1022, 1023, 1024, 1034
8051	39380042 (12193, 12194)	Novel Protein sim. GBank gl 416762 sp P32797 CC13_YEAST - CELL DIVISION CONTROL PROTEIN 13		UNCLASSIFIED	1004, 1022, 1024, 1034

8052	78725799 (12753, 12754)	Novel Protein sim. GBank gil416765 sp P25655 NOT1_YEAST - GENERAL NEGATIVE REGULATOR OF TRANSCRIPTION SUBUNIT 1		UNCLASSIFIED	1008, 1026
8053	28491387 (12973, 12974)	Novel Protein sim. GBank gil416765 sp P25655 NOT1_YEAST - GENERAL NEGATIVE REGULATOR OF TRANSCRIPTION SUBUNIT 1		UNCLASSIFIED	1008
8054	80052502 (9597, 9598)	Novel Protein sim. GBank gil416919 sp P32601 DSS4_YEAST - DSS4 PROTEIN		UNCLASSIFIED	1003, 1004, 1006, 1008, 1010, 1017, 1022, 1023, 1024, 1034, 1038
8055	8522425 (8441, 8442)	Novel Protein sim. GBank gil416935 sp P32324 EF2_YEAST - ELONGATION FACTOR 2 (EF-2)		UNCLASSIFIED	1010
8056	80051055 (413, 414)	Novel Protein sim. GBank gil417043 sp P32064 GCVA_ECOLI - GLYCINE CLEAVAGE SYSTEM TRANSCRIPTIONAL ACTIVATOR (GCV OPERON ACTIVATOR)		UNCLASSIFIED	1004, 1034
8057	16835402 (12339, 12340)	Novel Protein sim. GBank gil417117 sp P32828 HEX3_YEAST - HEXOSE METABOLISM-RELATED PROTEIN HEX3		UNCLASSIFIED	1017
8058	17707002 (397, 398)	Novel Protein sim. GBank gil417229 sp P32893 KR11_YEAST - BETA- GLUCAN SYNTHESIS-ASSOCIATED PROTEIN KRE11 (KILLER TOXIN-RESISTANCE PROTEIN 11)		UNCLASSIFIED	1017
8059	28790964 (19283, 19284)	Novel Protein sim. GBank gil417246 sp Q04760 LGUL_HUMAN - LACTOYLGLUTATHIONE LYASE (METHYLGLYOXALASE) (ALDOKETOMUTASE) (GLYOXALASE I) (GLX I) (KETONE-ALDEHYDE MUTASE) (S-D-LACTOYLGLUTATHIONE METHYLGLYOXAL LYASE)		UNCLASSIFIED	1022, 1026

8060	79258550 (10839, 10840)	Novel Protein sim. GBank gil417318 splP32335 MSS51_YEAST - MSS51 PROTEIN		UNCLASSIFIED	1003, 1022
8061	46661424 (6485, 6486)	Novel Protein sim. GBank gil417689 splP32843 RN12_YEAST - RNA12 PROTEIN		UNCLASSIFIED	1040
8062	17134806 (1621, 1622)	Novel Protein sim. GBank gil417726 splP32368 RSD1_YEAST - RECESSIVE SUPPRESSOR OF SECRETORY DEFECT		UNCLASSIFIED	1044
8063	79206539 (1457, 1458)	Novel Protein sim. GBank gil417752 splP32564 SCM4_YEAST - SCM4 PROTEIN		UNCLASSIFIED	1054
8064	80244127 (12079, 12080)	Novel Protein sim. GBank gil417774 splP32578 SIP1_YEAST - SIP1 PROTEIN		UNCLASSIFIED	1006, 1034
8065	78794590 (783, 784)	Novel Protein sim. GBank gil417782 splP32567 SMP2_YEAST - SMP2 PROTEIN		UNCLASSIFIED	1044
8066	25234413 (21023, 21024)	Novel Protein sim. GBank gil417782 splP32567 SMP2_YEAST - SMP2 PROTEIN		UNCLASSIFIED	1017, 1044
8067	30609808 (17697, 17698)	Novel Protein sim. GBank gil418427 splP32634 YEW2_YEAST - HYPOTHETICAL 195.4 KD PROTEIN IN RPS26B-GLC7 INTERGENIC REGION		UNCLASSIFIED	1022, 1026
8068	78795658 (21181, 21182)	Novel Protein sim. GBank gil418430 splP32645 ISC1_YEAST - MEIOSIS SPECIFIC PROTEIN ISC10		UNCLASSIFIED	1009
8069	20457900 (7899, 7900)	Novel Protein sim. GBank gil418436 splP32066 YGDE_ECOLI - HYPOTHETICAL 41.9 KD PROTEIN IN FUCR-GCVA INTERGENIC REGION (ORF3)		UNCLASSIFIED	1010
8070	27874349 (12897, 12898)	Novel Protein sim. GBank gil418501 splP32159 YIP_ECOLI - HYPOTHETICAL 32.9 KD PROTEIN IN CPXA-PFKA INTERGENIC REGION		UNCLASSIFIED	1022

8071	80230799 (5823, 5824)	Novel Protein sim. GBank gi 418609 sp P32464 YKS9_YEAST - HYPOTHETICAL 45.9 KD PROTEIN IN ONB1-PXA2 INTERGENIC REGION		UNCLASSIFIED	1001, 1006, 1008, 1009, 1010, 1017, 1022, 1024, 1034, 1038, 1039
8072	70336256 (1167, 1168)	Novel Protein sim. GBank gi 419916 pir JN0454 - hypothetical 10.5K protein (URA1 5' region) - gill mushroom (Agrocybe aegerita)		UNCLASSIFIED	1049
8073	79631076 (11647, 11648)	Novel Protein sim. GBank gi 4210557 dbj BAA74549 - (AB006372) maturase [Cotylelobium malayanum]		UNCLASSIFIED	1044
8074	87454411 (21823, 21624)	Novel Protein sim. GBank gi 4218120 emb CAA22974.1 - (AL035353) Proline-rich APG-like protein [Arabidopsis thaliana]		UNCLASSIFIED	1025
8075	28392349 (2293, 2294)	Novel Protein sim. GBank gi 4220529 emb CAA23002 - (AL035356) putative protein [Arabidopsis thaliana]		UNCLASSIFIED	1017
8076	77774744 (15211, 15212)	Novel Protein sim. GBank gi 423149 pir J46010 - X-linked retinopathy protein (C-terminal, clone XEH.8c) - human (fragment)		UNCLASSIFIED	1046
8077	13513214 (5805, 5806)	Novel Protein sim. GBank gi 4235480 gb AAD13223.1 - (AF100302) methyl-accepting chemotaxis protein [Pseudomonas putida]		UNCLASSIFIED	1024
8078	10047878 (18451, 18452)	Novel Protein sim. GBank gi 4235480 gb AAD13223.1 - (AF100302) methyl-accepting chemotaxis protein [Pseudomonas putida]		UNCLASSIFIED	1039
8079	86380109 (20823, 20824)	Novel Protein sim. GBank gi 423681 pir J533478 - hypothetical protein 2 - rat		UNCLASSIFIED	1053
8080	7653159 (21805, 21806)	Novel Protein sim. GBank gi 4240203 dbj BAA74880.1 - (AB020664) KIAA0857 protein [Homo sapiens]		UNCLASSIFIED	1034

8081	20731079 (14133, 14134)	Novel Protein sim. GBank gil4240241 dbj BAA74899.1 - (AB020683) KIAA0876 protein [Homo sapiens]		UNCLASSIFIED	1006, 1022
8082	94240626 (7089, 7090)	Novel Protein sim. GBank gil4249381 gb AAD14478 - (AC005966) Strong similarity to gb AF039376 Evelknievel retrotransposon polypeptide from Arabidopsis arenosa. [Arabidopsis thaliana]		UNCLASSIFIED	1017
8083	12659751 (3511, 3512)	Novel Protein sim. GBank gil4263825 gb AAD15468 - (AC006067) hypothetical protein [Arabidopsis thaliana]		UNCLASSIFIED	1004, 1006
8084	78769688 (5571, 5572)	Novel Protein sim. GBank gil4314370 gb AAD15581 - (AC006340) hypothetical protein [Arabidopsis thaliana]		UNCLASSIFIED	1022
8085	17301239 (9421, 9422)	Novel Protein sim. GBank gil4314387 gb AAD15597 - (AC006232) putative beta-alanine synthetase [Arabidopsis thaliana]		UNCLASSIFIED	1003
8086	29242476 (4107, 4108)	Novel Protein sim. GBank gil4320 emb CAA36300 - (X52081) RGP1 (reduced growth phenotype) gene product (AA 1- 661) [Saccharomyces cerevisiae]		UNCLASSIFIED	1017
8087	25324972 (9181, 9182)	Novel Protein sim. GBank gil4325123 gb AAD17271 - (AF119361) unknown [Frankia sp. EulK1]		UNCLASSIFIED	1003
8088	27962602 (13759, 13760)	Novel Protein sim. GBank gil4325354 gb AAD17351 - (AF128395) contains similarity to retrovirus-related polypeptides and to CCHC zinc finger protein (Pfam: PF00098, Score=16.3, E=0.051, E= 1) [Arabidopsis thaliana]		UNCLASSIFIED	1034
8089	29676045 (16353, 16354)	Novel Protein sim. GBank gil4328 emb CAA47121 - (X66501) RIF1 [Saccharomyces cerevisiae]		UNCLASSIFIED	1009
8090	57294469 (20567, 20568)	Novel Protein sim. GBank gil4335941 gb AAD17527 - (AF061738) leucine amino peptidase [Homo sapiens]		UNCLASSIFIED	1024

8091	80566418 (3743, 3744)	Novel Protein sim. GBank gi 4336418 gb AAD17796 - (AF086827) low voltage-activated T-type calcium channel alpha-1 subunit [Rattus norvegicus]		UNCLASSIFIED	1058
8092	80186916 (4483, 4484)	Novel Protein sim. GBank gi 4378025 gb AAD19348 - (AF086712) reverse transcriptase-like protein [Fugu rubripes]		UNCLASSIFIED	1058
8093	6626263 (18745, 18746)	Novel Protein sim. GBank gi 4378168 gb AAD19413.1 - (AF102543) unknown [Zymomonas mobilis]		UNCLASSIFIED	1022
8094	78405879 (1717, 1718)	Novel Protein sim. GBank gi 439877 L27428 - reverse transcriptase [Homo sapiens]		UNCLASSIFIED	1024
8095	65974969 (12085, 12086)	Novel Protein sim. GBank gi 439877 L27428 - reverse transcriptase [Homo sapiens]		UNCLASSIFIED	1046
8096	11357885 (15313, 15314)	Novel Protein sim. GBank gi 439877 L27428 - reverse transcriptase [Homo sapiens]		UNCLASSIFIED	1058
8097	27364200 (6891, 6892)	Novel Protein sim. GBank gi 4406513 gb AAD20010 - (AF123319) FtsI [Streptomyces coelicolor]		UNCLASSIFIED	1008
8098	78676958 (13057, 13058)	Novel Protein sim. GBank gi 4406581 gb AAD20037 - (AF131760) Unknown [Homo sapiens]		UNCLASSIFIED	1017
8099	56073773 (3439, 3440)	Novel Protein sim. GBank gi 4415912 gb AAD20143 - (AC006282) putative protease [Arabidopsis thaliana]		UNCLASSIFIED	1021
8100	10883621 (11541, 11542)	Novel Protein sim. GBank gi 4416479 gb AAD20379 - (AF125999) O-acetyl-transferase atfA [Mycobacterium avium]		UNCLASSIFIED	1024
8101	78526789 (17245, 17246)	Novel Protein sim. GBank gi 4417286 gb AAD20411 - (AC007019) putative shikimate kinase [Arabidopsis thaliana]		UNCLASSIFIED	1009
8102	80248339 (10893, 10894)	Novel Protein sim. GBank gi 4417309 gb AAD20433 - (AC006446) putative pol polyprotein with a reverse transcriptase domain [Arabidopsis thaliana]		UNCLASSIFIED	1006

8103	87448916 (22545, 22546)	Novel Protein sim. GBank gi 4432830 gb AAD20679 - (AC006283) similar to pheromone receptor deficient mutant [Arabidopsis thaliana]		UNCLASSIFIED	1007, 1025
8104	91235451 (21311, 21312)	Novel Protein sim. GBank gi 4432854 gb AAD20702 - (AC006300) hypothetical protein [Arabidopsis thaliana]		UNCLASSIFIED	1010
8105	65692088 (5381, 5382)	Novel Protein sim. GBank gi 4454023 emb CAA23020.1 - (AL035394) putative protein (fragment) [Arabidopsis thaliana]		UNCLASSIFIED	1009
8106	35106779 (21411, 21412)	Novel Protein sim. GBank gi 4455678 emb CAB36570 - (AL035472) putative protein synthesis initiation factor 2 [Mycobacterium leprae]		UNCLASSIFIED	1026, 1038
8107	27965088 (13249, 13250)	Novel Protein sim. GBank gi 4467274 emb CAB37598 - (AL035570) putative nitrilase homolog [Schizosaccharomyces pombe]		UNCLASSIFIED	1034
8108	78724598 (18621, 18622)	Novel Protein sim. GBank gi 4468887 emb CAB38141.1 - (AL035591) conserved hypothetical protein [Streptomyces coelicolor]		UNCLASSIFIED	1022
8109	37804149 (17945, 17946)	Novel Protein sim. GBank gi 4490702 emb CAB38836.1 - (AL035680) hypothetical protein [Arabidopsis thaliana]		UNCLASSIFIED	1058
8110	65897790 (5205, 5206)	Novel Protein sim. GBank gi 4493956 emb CAB1123.2 - (Z98551) predicted using hexExon; MAL3P6.28 (PFC0845c), Hypothetical protein, len: 167 aa; Similarity to model organism hypothetical proteins (C.elegans, D.melanogaster, S.cerevisiae & S.pombe). C.elegans protein ZK287.5 (TR:....		UNCLASSIFIED	1054
8111	98004396 (17447, 17448)	Novel Protein sim. GBank gi 4503037 ref NP_003842.1 pCREG - cellular repressor of E1A-stimulated genes		UNCLASSIFIED	1036, 1040, 1058

8112	25248298 (7757, 7758)	Novel Protein sim. GBank gi 4503181 ref NP_003919.1 pCXX1 - UNKNOWN		UNCLASSIFIED	1023
8113	78742048 (13949, 13950)	Novel Protein sim. GBank gi 4503355 ref NP_001371.1 pDOCK - dedicator of cyto-kinesis 1		UNCLASSIFIED	1022
8114	79212856 (11145, 11146)	Novel Protein sim. GBank gi 4503431 ref NP_003485.1 pDYSF - Dysferlin		UNCLASSIFIED	1054
8115	34078604 (2987, 2988)	Novel Protein sim. GBank gi 4504329 ref NP_003701.1 pHAI - hepatocyte growth factor activator inhibitor precursor		UNCLASSIFIED	1000
8116	5647220 (12491, 12492)	Novel Protein sim. GBank gi 4505891 ref NP_001075.1 pPLOC - UNKNOWN		UNCLASSIFIED	1058
8117	16845037 (11059, 11060)	Novel Protein sim. GBank gi 4506427 ref NP_002880.1 pRARR - retinoic acid receptor responder (tazarotene induced) 2		UNCLASSIFIED	1039
8118	11241746 (14555, 14556)	Novel Protein sim. GBank gi 4507053 ref NP_003974.1 pSLC7 - UNKNOWN		UNCLASSIFIED	1006
8119	5827976 (22455, 22456)	Novel Protein sim. GBank gi 4507367 ref NP_003182.1 pTARS - threonyl-tRNA synthetase		UNCLASSIFIED	1058
8120	91235061 (5551, 5552)	Novel Protein sim. GBank gi 4507529 ref NP_003255.1 pTLR2 - toll-like receptor2		UNCLASSIFIED	1010
8121	8362088 (15167, 15168)	Novel Protein sim. GBank gi 4507551 ref NP_003265.1 pTMEM - transmembrane protein 1		UNCLASSIFIED	1010
8122	80249007 (5209, 5210)	Novel Protein sim. GBank gi 4512375 db BAA75339.1 - (AB011837) phosphotransferase system (PTS) fructose-specific enzyme IIBC component [Bacillus halodurans]		UNCLASSIFIED	1006, 1010, 1023, 1053

8123	78676096 (11633, 11634)	Novel Protein sim. GBank gil4512375 dbj BAA75339.1 - (AB011837) phosphotransferase system (PTS) fructose- specific enzyme IBC component [Bacillus halodurans]		UNCLASSIFIED	1008, 1026
8124	87452028 (19503, 19504)	Novel Protein sim. GBank gil4512613 gb AAD21682.1 - (AC004793) F28K20.12 [Arabidopsis thaliana]		UNCLASSIFIED	1003
8125	87452033 (20495, 20496)	Novel Protein sim. GBank gil4512613 gb AAD21682.1 - (AC004793) F28K20.12 [Arabidopsis thaliana]		UNCLASSIFIED	1025
8126	11241968 (18377, 18378)	Novel Protein sim. GBank gil4512671 gb AAD21725.1 - (AC006931) unknown protein [Arabidopsis thaliana]		UNCLASSIFIED	1022
8127	30229992 (11587, 11588)	Novel Protein sim. GBank gil4512678 gb AAD21732.1 - (AC006931) unknown protein [Arabidopsis thaliana]		UNCLASSIFIED	1026
8128	87453856 (14531, 14532)	Novel Protein sim. GBank gil4512685 gb AAD21739.1 - (AC006931) hypothetical protein [Arabidopsis thaliana]		UNCLASSIFIED	1025
8129	78468948 (7957, 7958)	Novel Protein sim. GBank gil451544 (U04267) - proline-rich cell wall protein [Gossypium barbadense]		UNCLASSIFIED	1026
8130	79574476 (9003, 9004)	Novel Protein sim. GBank gil451544 (U04267) - proline-rich cell wall protein [Gossypium barbadense]		UNCLASSIFIED	1031
8131	23333928 (8777, 8778)	Novel Protein sim. GBank gil4538923 emb CAB39659.1 - (AL049483) predicted protein destination factor [Arabidopsis thaliana]		UNCLASSIFIED	1053
8132	71088499 (19799, 19800)	Novel Protein sim. GBank gil4539200 emb CAB39858.1 - (AL049497) putative long chain fatty acid coA ligase [Streptomyces coelicolor]		UNCLASSIFIED	1029, 1053
8133	11707058 (15811, 15812)	Novel Protein sim. GBank gil4539386 emb CAB37452.1 - (AL035526) extensin-like protein [Arabidopsis thaliana]		UNCLASSIFIED	1031

8134	79376717 (6883, 6884)	Novel Protein sim. GBank gil4557335[ref NP_000040.1 pASPA - aspartoacylase (aminoacylase 2)]		UNCLASSIFIED	1000, 1004, 1026, 1029
8135	65899415 (4583, 4584)	Novel Protein sim. GBank gil4559353[gb AAD23014.1 AC00658 - (AC006585) putative extragenic suppressor protein [Arabidopsis thaliana]]		UNCLASSIFIED	1044
8136	87453348 (13325, 13326)	Novel Protein sim. GBank gil4559384[gb AAD23044.1 AC00652 - (AC006526) unknown protein [Arabidopsis thaliana]]		UNCLASSIFIED	1025
8137	87452988 (15867, 15868)	Novel Protein sim. GBank gil4567281[gb AAD23694.1 AC00684 - (AC006841) unknown protein [Arabidopsis thaliana]]		UNCLASSIFIED	1008, 1025
8138	17890449 (5597, 5598)	Novel Protein sim. GBank gil4567305[gb AAD23716.1 AC00595 - (AC005956) hypothetical protein [Arabidopsis thaliana]]		UNCLASSIFIED	1044
8139	79574848 (14153, 14154)	Novel Protein sim. GBank gil4572367[gb AAD23762.1 AF11856 - (AF118562) evectin-1 [Rattus norvegicus]]		UNCLASSIFIED	1031
8140	28399367 (21963, 21964)	Novel Protein sim. GBank gil4574119[gb AAD23899.1 AF00941 - (AF009415) putative regulatory protein [Staphylococcus xyloso]]		UNCLASSIFIED	1023
8141	17900359 (3435, 3436)	Novel Protein sim. GBank gil4580331[emb CAB40107.1 - (AJ001206) putative glycogen debranching enzyme [Streptomyces coelicolor]]		UNCLASSIFIED	1044
8142	78792051 (4027, 4028)	Novel Protein sim. GBank gil4580997[gb AAD24571.1 AF12108 - (AF121081) cAMP inducible 2 protein [Mus musculus]]		UNCLASSIFIED	1017

8143	29352092 (7817, 7818)	Novel Protein sim. GBank gil4582390 emb CAB40337.1 - (AL049573) putative integral membrane efflux protein [Streptomyces coelicolor]		UNCLASSIFIED	1022
8144	28393172 (13691, 13692)	Novel Protein sim. GBank gil4582436 gb AAD24822.1 AC00719 - (AC007196) unknown protein [Arabidopsis thaliana]		UNCLASSIFIED	1044
8145	87463268 (5199, 5200)	Novel Protein sim. GBank gil4584255 emb CAB40577.1 - (Y18471) SINA1p [Vitis vinifera]		UNCLASSIFIED	1025
8146	3374453 (9045, 9046)	Novel Protein sim. GBank gil4584480 emb CAB40683.1 - (AL049587) putative transcriptional regulator [Streptomyces coelicolor]		UNCLASSIFIED	1012
8147	80056480 (17603, 17604)	Novel Protein sim. GBank gil4585976 gb AAD25612.1 AC00528 - (AC005287) Unknown protein [Arabidopsis thaliana]		UNCLASSIFIED	1006
8148	27366401 (15777, 15778)	Novel Protein sim. GBank gil4587562 gb AAD25793.1 AC00655 - (AC006550) Belongs to PF 00583 Acetyltransferase (GNAT) family. [Arabidopsis thaliana]		UNCLASSIFIED	1003, 1017
8149	25259946 (20267, 20268)	Novel Protein sim. GBank gil4587886 gb AAD25912.1 AF06930 - (AF069302) Orf6 [Pedococcus pentosaceus]		UNCLASSIFIED	1008
8150	82285798 (13493, 13494)	Novel Protein sim. GBank gil4589285 gb AAD26430.1 AF13515 - (AF135154) ferric alcaligin siderophore receptor [Bordetella pertussis]		UNCLASSIFIED	1016
8151	13886535 (6323, 6324)	Novel Protein sim. GBank gil4589478 dbj BAA76767.1 - (AB023140) KIAA0923 protein [Homo sapiens]		UNCLASSIFIED	1034

8152	87893247 (16901, 16902)	Novel Protein sim. GBank gil461488 sp P34106 ALA2_PANMI - ALANINE AMINOTRANSFERASE 2 (GPT) (GLUTAMIC--PYRUVIC TRANSAMINASE 2) (GLUTAMIC--ALANINE TRANSAMINASE 2) (ALAAT-2)		UNCLASSIFIED	1025
8153	79602385 (3121, 3122)	Novel Protein sim. GBank gil461532 sp P34727 ARF_AJECA - ADP-RIBOSYLATION FACTOR		UNCLASSIFIED	1003, 1006, 1016, 1017
8154	11072137 (3703, 3704)	Novel Protein sim. GBank gil461698 sp P33322 CBF5_YEAST - CENTROMERE/MICROTUBULE BINDING PROTEIN CBF5 (CENTROMERE-BINDING FACTOR 5) (NUCLEOLAR PROTEIN CBF5) (P64)		UNCLASSIFIED	1034
8155	80077624 (22413, 22414)	Novel Protein sim. GBank gil462020 sp P06616 ERA_ECOLI - GTP-BINDING PROTEIN ERA		UNCLASSIFIED	1026, 1034
8156	39565586 (4795, 4796)	Novel Protein sim. GBank gil462168 sp P33892 GCN1_YEAST - TRANSLATIONAL ACTIVATOR GCN1		UNCLASSIFIED	1004, 1006, 1010, 1022, 1024
8157	80497338 (9377, 9378)	Novel Protein sim. GBank gil462168 sp P33892 GCN1_YEAST - TRANSLATIONAL ACTIVATOR GCN1		UNCLASSIFIED	1004, 1006, 1008, 1009, 1010, 1012, 1017, 1022, 1024, 1026, 1039
8158	29229733 (22687, 22688)	Novel Protein sim. GBank gil462168 sp P33892 GCN1_YEAST - TRANSLATIONAL ACTIVATOR GCN1		UNCLASSIFIED	1017
8159	27359768 (17587, 17588)	Novel Protein sim. GBank gil462653 sp P35198 MTH1_YEAST - MTH1 PROTEIN		UNCLASSIFIED	1026

8160	78949231 (9513, 9514)	Novel Protein sim. GBank gi 4646206 gb AAD26879.1 AC00723 - (AC007230) Contains similarity to gb D13630 KIAA0005 gene from Homo sapiens. ESTs gb T45345, gb T21086, gb R90360, gb T20468, gb T45191 and gb A1100459 come from this gene. [Arabidopsis thaliana]		UNCLASSIFIED	1039
8161	30178353 (185, 186)	Novel Protein sim. GBank gi 464802 sp P35177 SPT7_YEAST - TRANSCRIPTIONAL ACTIVATOR SPT7		UNCLASSIFIED	1008
8162	80237091 (125, 126)	Novel Protein sim. GBank gi 464817 sp P34167 IF4B_YEAST - EUKARYOTIC TRANSLATION INITIATION FACTOR 4B (EIF-4B)		UNCLASSIFIED	1004, 1006, 1008, 1022
8163	11703163 (2445, 2446)	Novel Protein sim. GBank gi 465616 sp P33921 YEJL_ECOLI - HYPOTHETICAL 8.3 KD PROTEIN IN RPLY- PROL INTERGENIC REGION		UNCLASSIFIED	1038
8164	20467798 (19225, 19226)	Novel Protein sim. GBank gi 465689 sp P34234 YKS3_YEAST - HYPOTHETICAL 34.3 KD PROTEIN IN ORD1- FAS1 INTERGENIC REGION		UNCLASSIFIED	1022
8165	78051064 (16181, 16182)	Novel Protein sim. GBank gi 465716 sp P34250 YKK5_YEAST - HYPOTHETICAL 125.6 KD PROTEIN IN AAT1- GFA1 INTERGENIC REGION		UNCLASSIFIED	1008
8166	79621716 (13055, 13056)	Novel Protein sim. GBank gi 465721 sp P34252 YKK8_YEAST - HYPOTHETICAL 52.3 KD PROTEIN IN HAP4- AAT1 INTERGENIC REGION		UNCLASSIFIED	1006
8167	14997457 (17389, 17390)	Novel Protein sim. GBank gi 4678662 emb CAB41074.1 - (AL049645) putative large ATP-binding protein [Streptomyces coelicolor]		UNCLASSIFIED	1024

8168	17658858 (11833, 11834)	Novel Protein sim. GBank gil4680212[gb]AAD27575.1 AF11417 - (AF11417) hypothetical protein [Sorghum bicolor]		UNCLASSIFIED	1017
8169	17288824 (21897, 21898)	Novel Protein sim. GBank gil468277 (L31491) - ORF1; putative [pTOM9]		UNCLASSIFIED	1039
8170	80217675 (8951, 8952)	Novel Protein sim. GBank gil4688948[emb]CAB41506.1 - (AJ236881) SIT protein [Mus musculus]		UNCLASSIFIED	1004, 1012, 1022, 1026, 1031, 1034
8171	20378291 (5287, 5288)	Novel Protein sim. GBank gil4688953[emb]CAB41451.1 - (X84222) beta- succinyl CoA synthetase precursor [Neocallimastix frontalis]		UNCLASSIFIED	1004
8172	80058701 (1379, 1380)	Novel Protein sim. GBank gil4753853[emb]CAB42016.1 - (AL049754) putative adenylosuccinate synthetase [Streptomyces coelicolor]		UNCLASSIFIED	1010
8173	66314831 (18009, 18010)	Novel Protein sim. GBank gil4757964[ref]NP_004056.1 pCDR1 - cerebellar degeneration-related protein (34kD)		UNCLASSIFIED	1016
8174	20689941 (10125, 10126)	Novel Protein sim. GBank gil4758602[ref]NP_004508.1 pILF2 - interleukin enhancer binding factor 2, 45kD		UNCLASSIFIED	1010
8175	21433872 (17771, 17772)	Novel Protein sim. GBank gil4760712[dbj]BAA77407.1 - (AB016984) Sperizin [Mus musculus]		UNCLASSIFIED	1004, 1022
8176	66410974 (1625, 1626)	Novel Protein sim. GBank gil4760782[dbj]BAA77399.1 - (AB025413) Ten- m4 [Mus musculus]		UNCLASSIFIED	1038
8177	40385052 (15567, 15568)	Novel Protein sim. GBank gil4760782[dbj]BAA77399.1 - (AB025413) Ten- m4 [Mus musculus]		UNCLASSIFIED	1037
8178	78104938 (2635, 2636)	Novel Protein sim. GBank gil4768836[gb]AAD29636.1 AF11690 - (AF11690) unknown [Homo sapiens]		UNCLASSIFIED	1040

8179	80499684 (14195, 14196)	Novel Protein sim. GBank gil4768865[gb]/AAD29659.1 AF12475 - (AF124757) 1-deoxy-D-xylose 5-phosphate reductoisomerase [Zymomonas mobilis]		UNCLASSIFIED	1012
8180	78792187 (10217, 10218)	Novel Protein sim. GBank gil477443[pir]/A49013 - HTS1 - human (fragment)		UNCLASSIFIED	1003, 1044
8181	19515552 (10147, 10148)	Novel Protein sim. GBank gil478673[pir]/S23737 - proline-rich protein precursor - kidney bean		UNCLASSIFIED	1010
8182	95295782 (8399, 8400)	Novel Protein sim. GBank gil4790[emb]/CAA49299] - (X69584) YKL253 [Saccharomyces cerevisiae]		UNCLASSIFIED	1001
8183	78928624 (3177, 3178)	Novel Protein sim. GBank gil4808945[gb]/AAD30027.1 AF11917 - (AF119170) unknown [Escherichia coli]		UNCLASSIFIED	1044
8184	11759530 (18719, 18720)	Novel Protein sim. GBank gil4826478[emb]/CAB42896.1] - (Z83844) similar to SH3-binding protein [Homo sapiens]		UNCLASSIFIED	1024
8185	80079498 (2963, 2964)	Novel Protein sim. GBank gil4835305[emb]/CAB42927.1] - (AL049863) putative dehydrogenase [Streptomyces coelicolor]		UNCLASSIFIED	1034
8186	23302170 (16199, 16200)	Novel Protein sim. GBank gil4836835[gb]/AAC96549.2] - (U42580) PBCV-1 chitinase [Paramoecium bursaria Chlorella virus 1]		UNCLASSIFIED	1023
8187	11076707 (10583, 10584)	Novel Protein sim. GBank gil4838477[gb]/AAD31012.1 AF12722 - (AF127222) outer membrane heme receptor PfrR [Pseudomonas fluorescens]		UNCLASSIFIED	1010
8188	71220821 (4189, 4190)	Novel Protein sim. GBank gil4868437[gb]/AAD31316.1 AF14405 - (AF144054) apoptosis related protein APR-4 [Homo sapiens]		UNCLASSIFIED	1024, 1044

8189	46763899 (21239, 21240)	Novel Protein sim. GBank gil488437 gb AAD31316.1 AF14405 - (AF144054) apoptosis related protein APR-4 [Homo sapiens]		UNCLASSIFIED	1033
8190	86462959 (14559, 14560)	Novel Protein sim. GBank gil4883453 emb CAB43167.1 - (AL049913) putative histidinol-phosphate aminotransferase [Mycobacterium leprae]		UNCLASSIFIED	1048
8191	78460334 (21735, 21736)	Novel Protein sim. GBank gil4883930 gb AAD31711.1 AF13432 - (AF134322) chimeric antifreeze glycoprotein- serine protease [Dissostichus mawsoni]		UNCLASSIFIED	1003
8192	78795108 (21851, 21852)	Novel Protein sim. GBank gil4884372 emb CAB43300.1 - (AL050159) hypothetical protein [Homo sapiens]		UNCLASSIFIED	1026
8193	65888815 (16959, 16960)	Novel Protein sim. GBank gil4885649 ref NP_005490.1 pUBA2 - SUMO-1 activating enzyme subunit 2		UNCLASSIFIED	1054
8194	28796999 (17727, 17728)	Novel Protein sim. GBank gil4886900 gb AAD32094.1 AF11761 - (AF117610) inner centromere protein INCENP [Mus musculus]		UNCLASSIFIED	1022
8195	7419649 (14375, 14376)	Novel Protein sim. GBank gil4887204 gb AAD32231.1 AF14744 - (AF147448) rod-shape-determining protein [Pseudomonas aeruginosa]		UNCLASSIFIED	1058
8196	13085947 (303, 304)	Novel Protein sim. GBank gil4887209 gb AAD32236.1 AF14744 - (AF147448) lipote biosynthesis protein B [Pseudomonas aeruginosa]		UNCLASSIFIED	1027
8197	79834598 (18661, 18662)	Novel Protein sim. GBank gil4887211 gb AAD32237.1 AF14744 - (AF147449) penicillin-binding protein 1B [Pseudomonas aeruginosa]		UNCLASSIFIED	1023, 1038
8198	80227151 (9289, 9290)	Novel Protein sim. GBank gil4895141 gb AAD32748.1 - (AF127374) MmcW [Streptomyces lavendulae]		UNCLASSIFIED	1008, 1009, 1010, 1016, 1022, 1024, 1039

8199	25153220 (11697, 11698)	Novel Protein sim. GBank gil4914402 emb CAB43653.1 - (AL050352) hypothetical protein [Arabidopsis thaliana]		UNCLASSIFIED	1009
8200	54994470 (12895, 12896)	Novel Protein sim. GBank gil4926959 gb AAD32966.1 AF10387 - (AF10387) transcription repressor [Erwinia carotovora subsp. carotovora]		UNCLASSIFIED	1029
8201	20296536 (15737, 15738)	Novel Protein sim. GBank gil4929540 gb AAD34030.1 AF15138 - (AF15138) S-adenosylmethionine decarboxylase [Neurospora crassa]		UNCLASSIFIED	1034
8202	17710289 (3693, 3694)	Novel Protein sim. GBank gil4929765 gb AAD34143.1 AF15190 - (AF15190) CGI-148 protein [Homo sapiens]		UNCLASSIFIED	1044
8203	17940130 (13261, 13262)	Novel Protein sim. GBank gil4972787 gb AAD34790.1 AF13640 - (AF13640) threonyl-tRNA synthetase [Pseudomonas fluorescens]		UNCLASSIFIED	1039
8204	8490247 (18709, 18710)	Novel Protein sim. GBank gil497619 (L25608) - N- acetyltransferase [Saccharomyces cerevisiae]		UNCLASSIFIED	1022
8205	86465057 (12041, 12042)	Novel Protein sim. GBank gil4980524 gb AAD35133.1 AE00169 - (AE00169) conserved hypothetical protein [Thermotoga maritima]		UNCLASSIFIED	1029
8206	27830677 (1689, 1690)	Novel Protein sim. GBank gil4980613 gb AAD35215.1 AE00169 - (AE00169) conserved hypothetical protein [Thermotoga maritima]		UNCLASSIFIED	1004
8207	79820158 (17431, 17432)	Novel Protein sim. GBank gil4980663 gb AAD35261.1 AE00170 - (AE00170) leucyl-tRNA synthetase [Thermotoga maritima]		UNCLASSIFIED	1006, 1008
8208	47656572 (19991, 19992)	Novel Protein sim. GBank gil4981025 gb AAD35598.1 AE00172 - (AE00172) comM protein [Thermotoga maritima]		UNCLASSIFIED	1029

8209	20294813 (20309, 20310)	Novel Protein sim. GBank gil4981266 gb AAD35822.1 AE00174 - (AE001744) lipopolysaccharide core biosynthesis protein KdtB [Thermotoga maritima]		UNCLASSIFIED	1034
8210	11103031 (10987, 10988)	Novel Protein sim. GBank gil4981964 gb AAD36471.1 AE00179 - (AE001793) aspartate aminotransferase, putative [Thermotoga maritima]		UNCLASSIFIED	1006
8211	80060366 (13945, 13946)	Novel Protein sim. GBank gil4982168 gb AAD36664.1 AE00180 - (AE001804) hypothetical protein [Thermotoga maritima]		UNCLASSIFIED	1010, 1024
8212	23292149 (18043, 18044)	Novel Protein sim. GBank gil4982442 gb AAD36919.1 AE00182 - (AE001823) conserved hypothetical protein [Thermotoga maritima]		UNCLASSIFIED	1026
8213	9862603 (21667, 21668)	Novel Protein sim. GBank gil498253 (U02372) - integrase [Vibrio cholerae]		UNCLASSIFIED	1008
8214	78968895 (8963, 8964)	Novel Protein sim. GBank gil4995890 emb CAB44316.1 - (AJ242659) serine palmitoyltransferase [Solanum tuberosum]		UNCLASSIFIED	1039
8215	19526575 (4163, 4164)	Novel Protein sim. GBank gil4995986 dbj BAA78219.1 - (AB021506) 13.6% identical to DR8 gene of strain U1102 of HHV-6 [Human herpesvirus 6]		UNCLASSIFIED	1034
8216	20751378 (6411, 6412)	Novel Protein sim. GBank gil4996501 dbj BAA78502.1 - (AB021827) serine repeat antigen [Plasmodium falciparum]		UNCLASSIFIED	1022
8217	7660530 (12639, 12640)	Novel Protein sim. GBank gil5006991 gb AAD37772.1 AF14653 - (AF146532) glucosyl transferase WaaE [Klebsiella pneumoniae]		UNCLASSIFIED	1034
8218	80588367 (12431, 12432)	Novel Protein sim. GBank gil5031275 gb AAD38143.1 AF13949 - (AF139496) unknown [Prunus armeniaca]		UNCLASSIFIED	1053, 1058

8219	28467806 (21951, 21952)	Novel Protein sim. GBank gil5031775 ref NP_005799.1 pHYA2 - HYA22 protein		UNCLASSIFIED	1008
8220	57299447 (22393, 22394)	Novel Protein sim. GBank gil5031793 ref NP_005528.1 pING1 - inhibitor of growth 1		UNCLASSIFIED	1004, 1010, 1022, 1034, 1038
8221	86468627 (10411, 10412)	Novel Protein sim. GBank gil5031889 ref NP_005570.1 pLRE1 - LINE retrotransposable element 1		UNCLASSIFIED	1029
8222	78727868 (8703, 8704)	Novel Protein sim. GBank gil5032183 ref NP_005808.1 pTIP4 - cargo selection protein (mannose 6 phosphate receptor binding protein)		UNCLASSIFIED	1022
8223	7511820 (1815, 1816)	Novel Protein sim. GBank gil5032247 ref NP_005788.1 pEVA1 - epithelial V-like antigen		UNCLASSIFIED	1058
8224	23308583 (22591, 22592)	Novel Protein sim. GBank gil5042155 emb CAB44674.1 - (AL078620) putative protein [Arabidopsis thaliana]		UNCLASSIFIED	1053
8225	94233781 (16459, 16460)	Novel Protein sim. GBank gil5042457 gb AAD38294.1 AC00778 - (AC007789) unknown protein [Oryza sativa]		UNCLASSIFIED	1003
8226	78911515 (16031, 16032)	Novel Protein sim. GBank gil504480 (U10470) - unknown [Pseudomonas fluorescens]		UNCLASSIFIED	1003
8227	85822661 (5919, 5920)	Novel Protein sim. GBank gil5059250 gb AAD38955.1 - (AF147978) putative portal protein [Bacteriophage D3]		UNCLASSIFIED	1019
8228	87452751 (5935, 5936)	Novel Protein sim. GBank gil5081610 gb AAD39464.1 AF13544 - (AF135440) huntington yeast partner C [Mus musculus]		UNCLASSIFIED	1025
8229	78527630 (12341, 12342)	Novel Protein sim. GBank gil5091509 dbj BAA78744.1 - (AB023482) EST AU065533 (C2174) corresponds to a region of the predicted gene. Similar to Homo sapiens splicing factor Prp8 mRNA, complete cds. (AF092565) [Oryza sativa]		UNCLASSIFIED	1003

8230	87913676 (20147, 20148)	Novel Protein sim. GBank gil5091509 dbj BAA78744.1 - (AB023482) EST AU065533(C2174) corresponds to a region of the predicted gene.; Similar to Homo sapiens splicing factor Prip8 mRNA, complete cds. (AF092565) [Oryza sativa]		UNCLASSIFIED	1025
8231	56701283 (9209, 9210)	Novel Protein sim. GBank gil5102785 emb CAB45200.1 - (AL079308) putative transcriptional regulator [Streptomyces coelicolor]		UNCLASSIFIED	1010
8232	9403895 (21193, 21194)	Novel Protein sim. GBank gil5103613 dbj BAA79134.1 - (AP000058) 318aa long hypothetical carboxyphosphoenolpyruvate phosphonmutase [Aeropyrum pernix]		UNCLASSIFIED	1044
8233	11398393 (15621, 15622)	Novel Protein sim. GBank gil5104764 dbj BAA80079.1 - (AP000060) 619aa long hypothetical DNA ligase [Aeropyrum pernix]		UNCLASSIFIED	1034
8234	79574341 (2671, 2672)	Novel Protein sim. GBank gil5105920 dbj BAA81232.1 - (AP000063) 297aa long hypothetical protein [Aeropyrum pernix]		UNCLASSIFIED	1031
8235	9203662 (947, 948)	Novel Protein sim. GBank gil5106233 dbj BAA81544.1 - (AP000064) 687aa long hypothetical hydantoinase [Aeropyrum pernix]		UNCLASSIFIED	1023
8236	95086920 (21809, 21810)	Novel Protein sim. GBank gil5106572 gb AAD39760.1 AF14394 - (AF143946) transcriptional activator SRCAP [Homo sapiens]		UNCLASSIFIED	1003, 1024, 1030, 1038
8237	11103612 (597, 598)	Novel Protein sim. GBank gil520476 (U09129) - Mkt1p [Saccharomyces cerevisiae]		UNCLASSIFIED	1006, 1022
8238	46731718 (20527, 20528)	Novel Protein sim. GBank gil5257454 gb AAD41342.1 AF14432 - (AF144325) heparanase [Homo sapiens]		UNCLASSIFIED	1037

8239	80084413 (19763, 19764)	Novel Protein sim. GBank gil5306245 gb AAD41978.1 AC00643 - (AC006438) unknown protein [Arabidopsis thaliana]		UNCLASSIFIED	1001, 1022, 1034
8240	91216889 (5755, 5756)	Novel Protein sim. GBank gil5360958 emb CAA76860.2 - (Y17797) hypothetical protein [Enterococcus faecalis]		UNCLASSIFIED	1004
8241	47656880 (5951, 5952)	Novel Protein sim. GBank gil539513 pir A47718 - reverse transcriptase, pol-like - chicken (fragment)		UNCLASSIFIED	1029
8242	80062436 (15263, 15264)	Novel Protein sim. GBank gil541168 pir S40860 - periplasmic sulphate binding protein - Escherichia coli		UNCLASSIFIED	1010
8243	71817777 (17931, 17932)	Novel Protein sim. GBank gil5419988 emb CAB46500.1 - (AJ243503) m46 protein [Mus musculus]		UNCLASSIFIED	1008
8244	80504136 (2903, 2904)	Novel Protein sim. GBank gil5420387 emb CAB46679.1 - (AJ243459) proteophosphoglycan [Leishmania major]		UNCLASSIFIED	1012
8245	80049411 (4725, 4726)	Novel Protein sim. GBank gil5420387 emb CAB46679.1 - (AJ243459) proteophosphoglycan [Leishmania major]		UNCLASSIFIED	1010
8246	20465367 (7559, 7560)	Novel Protein sim. GBank gil5420387 emb CAB46679.1 - (AJ243459) proteophosphoglycan [Leishmania major]		UNCLASSIFIED	1010
8247	80497336 (9195, 9196)	Novel Protein sim. GBank gil5420387 emb CAB46679.1 - (AJ243459) proteophosphoglycan [Leishmania major]		UNCLASSIFIED	1012
8248	79483267 (9469, 9470)	Novel Protein sim. GBank gil5420387 emb CAB46679.1 - (AJ243459) proteophosphoglycan [Leishmania major]		UNCLASSIFIED	1006, 1027
8249	20158809 (10629, 10630)	Novel Protein sim. GBank gil5420387 emb CAB46679.1 - (AJ243459) proteophosphoglycan [Leishmania major]		UNCLASSIFIED	1010
8250	100328608 (12445, 12446)	Novel Protein sim. GBank gil5420387 emb CAB46679.1 - (AJ243459) proteophosphoglycan [Leishmania major]		UNCLASSIFIED	1017, 1024, 1038, 1039, 1044

8251	19748016 (19611, 19612)	Novel Protein sim. GBank gil5420387[emb CAB46679.1] - (AJ243459) proteophosphoglycan [Leishmania major]		UNCLASSIFIED	1003
8252	78707671 (5861, 5862)	Novel Protein sim. GBank gil5430762[gb AAD43162.1 AC00750 - (AC007504) Similar to CCS1 [Arabidopsis thaliana]		UNCLASSIFIED	1025
8253	8210021 (2155, 2156)	Novel Protein sim. GBank gil5453573[ref NP_006411.1 pBIG2 - brefeldin A- inhibited guanine nucleotide-exchange protein 2		UNCLASSIFIED	1034
8254	86380228 (67, 68)	Novel Protein sim. GBank gil5453623[ref NP_006305.1 pCNK1 - connector enhancer of KSR-like (Drosophila kinase suppressor of ras)		UNCLASSIFIED	1053
8255	95357725 (2985, 2986)	Novel Protein sim. GBank gil5453702[ref NP_006294.1 pJTV1 - JTV1 gene		UNCLASSIFIED	1054
8256	24136710 (2329, 2330)	Novel Protein sim. GBank gil5457277[emb CAB46965.1] - (AL096823) putative membrane protein [Streptomyces coelicolor]		UNCLASSIFIED	1003
8257	20482246 (7935, 7936)	Novel Protein sim. GBank gil5457625[emb CAB49116.1] - (AJ248283) PAB2227 [Pyrococcus abyssi]		UNCLASSIFIED	1010
8258	29346708 (16723, 16724)	Novel Protein sim. GBank gil547573[emb CAA79376] - (Z18944) ORF1 [Saccharomyces cerevisiae]		UNCLASSIFIED	1022
8259	88097318 (14895, 14896)	Novel Protein sim. GBank gil547648[sp P36605 HIS8_SCHPO - HISTIDINOL PHOSPHATE AMINOTRANSFERASE (IMIDAZOLE ACETOL-PHOSPHATE TRANSAMINASE)		UNCLASSIFIED	1018
8260	20724456 (19731, 19732)	Novel Protein sim. GBank gil547653[sp P35915 HMGL_CHICK - HYDROXYMETHYLGLUTARYL-COA LYASE (HMG-COA LYASE) (HL) (3-HYDROXY-3- METHYLGLUTARATE-COA LYASE)		UNCLASSIFIED	1006

8261	78527714 (10789, 10790)	Novel Protein sim. GBank gil549010 sp P35614 ERF1_ARATH - EUKARYOTIC PEPTIDE CHAIN RELEASE FACTOR SUBUNIT 1 (ERF1) (OMNIPOTENT SUPPRESSOR PROTEIN 1 HOMOLOG) (SUP1 HOMOLOG)		UNCLASSIFIED	1003
8262	37036342 (16023, 16024)	Novel Protein sim. GBank gil549100 sp Q04222 TRA1_ALCSP - TRANSPOSASE FOR INSERTION SEQUENCE ELEMENT IS1071 IN TRANSPOSON TN5271		UNCLASSIFIED	1012
8263	79185719 (13639, 13640)	Novel Protein sim. GBank gil549542 sp P36999 RRMA_ECOLI - RRNA (GUANINE-N1-J)-METHYLTRANSFERASE (23S RRNA M1G745 METHYLTRANSFERASE)		UNCLASSIFIED	1022
8264	20728483 (8365, 8366)	Novel Protein sim. GBank gil549560 sp P37013 YGAA_ECOLI - HYPOTHETICAL SIGMA-54-DEPENDENT TRANSCRIPTIONAL REGULATOR IN GUTQ- HYPF INTERGENIC REGION		UNCLASSIFIED	1006
8265	78384675 (12307, 12308)	Novel Protein sim. GBank gil549592 sp P36246 RIMM_SALTY - 16S RRNA PROCESSING PROTEIN RIMM		UNCLASSIFIED	1026
8266	80219688 (15817, 15818)	Novel Protein sim. GBank gil549598 sp P36121 YK05_YEAST - HYPOTHETICAL 32.1 KD PROTEIN IN DBP7- GCN3 INTERGENIC REGION		UNCLASSIFIED	1004, 1006, 1010, 1022, 1023, 1024, 1034
8267	79818407 (18311, 18312)	Novel Protein sim. GBank gil549635 sp P36157 YK57_YEAST - HYPOTHETICAL 39.9 KD PROTEIN IN SIS2- MTD1 INTERGENIC REGION		UNCLASSIFIED	1006, 1010, 1022, 1024
8268	30454255 (6409, 6410)	Novel Protein sim. GBank gil549637 sp P36159 YK59_YEAST - HYPOTHETICAL 96.8 KD PROTEIN IN SIS2- MTD1 INTERGENIC REGION		UNCLASSIFIED	1008

8269	11758274 (8619, 8620)	Novel Protein sim. GBank gil549646 sp P36168 YK76_YEAST - HYPOTHETICAL 137.5 KD PROTEIN IN MPL1- PPC1 INTERGENIC REGION		UNCLASSIFIED	1022
8270	20455715 (6617, 6618)	Novel Protein sim. GBank gil549652 sp P36173 YK86_YEAST - HYPOTHETICAL 69.0 KD PROTEIN IN SIR1 3'REGION		UNCLASSIFIED	1010, 1024
8271	83205017 (13267, 13268)	Novel Protein sim. GBank gil549663 sp P36101 YK7_YEAST - HYPOTHETICAL 50.3 KD PROTEIN IN TFA1- PAN3 INTERGENIC REGION		UNCLASSIFIED	1003, 1008, 1017, 1023, 1039, 1044
8272	28972283 (15385, 15386)	Novel Protein sim. GBank gil549675 sp P36090 YK7_YEAST - HYPOTHETICAL 58.9 KD PROTEIN IN ELM1- PRI2 INTERGENIC REGION		UNCLASSIFIED	1003
8273	38297099 (20321, 20322)	Novel Protein sim. GBank gil549730 sp P36055 YKP8_YEAST - HYPOTHETICAL 11.4 KD PROTEIN IN PIR3- APE2 INTERGENIC REGION		UNCLASSIFIED	1029
8274	79831949 (4647, 4648)	Novel Protein sim. GBank gil549767 sp P36111 YKZ5_YEAST - HYPOTHETICAL 66.6 KD PROTEIN IN YPT52- DBP7 INTERGENIC REGION		UNCLASSIFIED	1008
8275	79869439 (19073, 19074)	Novel Protein sim. GBank gil549767 sp P36111 YKZ5_YEAST - HYPOTHETICAL 66.6 KD PROTEIN IN YPT52- DBP7 INTERGENIC REGION		UNCLASSIFIED	1006, 1012
8276	26305361 (2551, 2552)	Novel Protein sim. GBank gil549770 sp P36114 YKZ8_YEAST - HYPOTHETICAL 81.8 KD PROTEIN IN YPT52- DBP7 INTERGENIC REGION		UNCLASSIFIED	1017
8277	78961607 (11613, 11614)	Novel Protein sim. GBank gil549771 sp P36115 YKZ9_YEAST - HYPOTHETICAL 68.9 KD PROTEIN IN YPT52- DBP7 INTERGENIC REGION		UNCLASSIFIED	1044

8278	30652671 (17325, 17326)	Novel Protein sim. GBank gil549806 spl Q06373 YPLC_CLOPE - HYPOTHETICAL 55.7 KD PROTEIN IN PLC 5'REGION (ORF 2)		UNCLASSIFIED	1022
8279	27931479 (5331, 5332)	Novel Protein sim. GBank gil550095 dbj BAA07427 - (D38310) Boi2p [Saccharomyces cerevisiae]		UNCLASSIFIED	1006
8280	35997569 (16795, 16796)	Novel Protein sim. GBank gil550426 emb CAA57411 - (X81813) CDC1 [Saccharomyces cerevisiae]		UNCLASSIFIED	1029
8281	20536647 (17989, 17990)	Novel Protein sim. GBank gil5521 emb CAA48518.1 - (X68506) GTPase- activating protein of Ypt6 [Saccharomyces cerevisiae]		UNCLASSIFIED	1058
8282	5730485 (15607, 15608)	Novel Protein sim. GBank gil5531829 gb AAD44489.1 - (AF078857) PTD002 [Homo sapiens]		UNCLASSIFIED	1058
8283	56946031 (12017, 12018)	Novel Protein sim. GBank gil5531843 gb AAD44496.1 - (AF078864) PTD011 [Homo sapiens]		UNCLASSIFIED	1001
8284	37797548 (14971, 14972)	Novel Protein sim. GBank gil5565917 gb AAD45266.1 AF14779 - (AF147795) WbjC [Pseudomonas aeruginosa]		UNCLASSIFIED	1012
8285	8490041 (18203, 18204)	Novel Protein sim. GBank gil556853 emb CAA86077 - (Z37896) incomplete orf, len: 744, CAI: 0.14 [Saccharomyces cerevisiae]		UNCLASSIFIED	1022
8286	21637724 (19541, 19542)	Novel Protein sim. GBank gil5578704 gb AAD17380.2 - (U65001) unknown [Sphingomonas sp.]		UNCLASSIFIED	1017
8287	7419799 (9683, 9684)	Novel Protein sim. GBank gil5579033 emb CAB51371.1 - (AJ249084) transposase [Pleuronectes platessa]		UNCLASSIFIED	1058
8288	25340671 (15387, 15388)	Novel Protein sim. GBank gil558251 emb CAA57607 - (X82086) D4446 [Saccharomyces cerevisiae]		UNCLASSIFIED	1003

8289	28848534 (16445, 16446)	Novel Protein sim. GBank gij5668607 gb AAD45978.1 - (AF115334) EstX [Pseudomonas fluorescens]		UNCLASSIFIED	1026
8290	36733824 (19721, 19722)	Novel Protein sim. GBank gij5689519 dbj BAA83043.1 - (AB029014) KIAA1091 protein [Homo sapiens]		UNCLASSIFIED	1026
8291	52415482 (4497, 4498)	Novel Protein sim. GBank gij5689890 emb CAB52053.1 - (AL109732) hypothetical protein [Streptomyces coelicolor A3(2)]		UNCLASSIFIED	1004, 1041
8292	94651627 (13363, 13364)	Novel Protein sim. GBank gij5689948 emb CAB51985.1 - (AL109663) putative isoleucyl-tRNA synthetase [Streptomyces coelicolor A3(2)]		UNCLASSIFIED	1010, 1022, 1024
8293	10311821 (21225, 21226)	Novel Protein sim. GBank gij5701961 emb CAB52153.1 - (AL109736) pelota protein/yeast dom34 homolog; probable involvement in meiotic and mitotic divisions [Schizosaccharomyces pombe]		UNCLASSIFIED	1001
8294	47652284 (10041, 10042)	Novel Protein sim. GBank gij5706510 emb CAB52272.1 - (AL109739) putative yeast shp1/ rat p47 homolog; contains UBX domain [Schizosaccharomyces pombe]		UNCLASSIFIED	1029
8295	80218138 (19501, 19502)	Novel Protein sim. GBank gij5726647 gb AAD48499.1 AF17368 - (AF173681) thioredoxin interacting factor [Mus musculus]		UNCLASSIFIED	1022
8296	80237351 (19723, 19724)	Novel Protein sim. GBank gij57806 emb CAA33483 - (X15443) gamma- glutamyltranspeptidase (AA 1-568) [Rattus sp.]		UNCLASSIFIED	1001, 1004, 1006, 1008, 1010, 1017, 1022, 1024, 1034
8297	7541650 (11623, 11624)	Novel Protein sim. GBank gij584850 sp P38041 BOB1_YEAST - BOB1 PROTEIN (BEM1-BINDING PROTEIN)		UNCLASSIFIED	1058

8298	78761583 (14347, 14348)	Novel Protein sim. GBank gil585255 sp P33734 HIS5_YEAST - HISTIDINE BIOSYNTHESIS BIFUNCTIONAL AMIDOTRANSFERASE / CYCLASE		UNCLASSIFIED	1017, 1038
8299	11611086 (17291, 17292)	Novel Protein sim. GBank gil585377 sp P38131 KTR4_YEAST - PROBABLE MANNOSYLTRANSFERASE KTR4		UNCLASSIFIED	1006
8300	80019584 (37, 38)	Novel Protein sim. GBank gil585473 sp P37958 MECA_BACSU - NEGATIVE REGULATOR OF GENETIC COMPETENCE MECA		UNCLASSIFIED	1006
8301	11815990 (6777, 6778)	Novel Protein sim. GBank gil585735 sp Q08423 PS2_MOUSE - PS2 PROTEIN PRECURSOR		UNCLASSIFIED	1006
8302	25260692 (6915, 6916)	Novel Protein sim. GBank gil585986 sp P38376 SECY_BREFL - PREPROTEIN TRANSLOCASE SECY SUBUNIT		UNCLASSIFIED	1008
8303	24145008 (17049, 17050)	Novel Protein sim. GBank gil586056 sp P00961 SYGB_ECOLI - GLYCYL-TRNA SYNTHETASE BETA CHAIN (GLYCINE--TRNA LIGASE BETA CHAIN) (GLYRS)		UNCLASSIFIED	1003
8304	66398087 (1703, 1704)	Novel Protein sim. GBank gil586292 sp P38118 YARF_CORGL - HYPOTHETICAL 26.1 KD PROTEIN IN AROF 3'REGION		UNCLASSIFIED	1009
8305	5706139 (1305, 1306)	Novel Protein sim. GBank gil586293 sp P38114 YB00_YEAST - PUTATIVE 126.9 KD TRANSCRIPTIONAL REGULATORY PROTEIN IN YSW1-RIB7 INTERGENIC REGION		UNCLASSIFIED	1058
8306	78903891 (195, 196)	Novel Protein sim. GBank gil586300 sp P38287 YB11_YEAST - HYPOTHETICAL 44.4 KD PROTEIN IN CDC28-ARL1 INTERGENIC REGION		UNCLASSIFIED	1003, 1008, 1022, 1044

8307	8362378 (3807, 3808)	Novel Protein sim. GBank gil586309 sp P38295 YB27_YEAST - HYPOTHETICAL 51.3 KD PROTEIN IN SMY2- RPS101 INTERGENIC REGION		UNCLASSIFIED	1004, 1058
8308	29358048 (3201, 3202)	Novel Protein sim. GBank gil586342 sp P38321 YB75_YEAST - HYPOTHETICAL 101.2 KD PROTEIN IN PCS60- ABD1 INTERGENIC REGION		UNCLASSIFIED	1009
8309	11613292 (3419, 3420)	Novel Protein sim. GBank gil586342 sp P38321 YB75_YEAST - HYPOTHETICAL 101.2 KD PROTEIN IN PCS60- ABD1 INTERGENIC REGION		UNCLASSIFIED	1010
8310	80219453 (2781, 2782)	Novel Protein sim. GBank gil586357 sp P38153 APM3_YEAST - ADAPTIN MEDIUM CHAIN HOMOLOG APM3		UNCLASSIFIED	1006, 1022, 1026, 1034
8311	79629266 (11191, 11192)	Novel Protein sim. GBank gil586361 sp P38358 YB8G_YEAST - HYPOTHETICAL 51.7 KD PROTEIN IN CTP1- SUL2 INTERGENIC REGION		UNCLASSIFIED	1025
8312	27838182 (7449, 7450)	Novel Protein sim. GBank gil586387 sp P38347 YB9P_YEAST - HYPOTHETICAL 48.0 KD PROTEIN IN MRPL37- RIF1 INTERGENIC REGION		UNCLASSIFIED	1006
8313	78794517 (19155, 19156)	Novel Protein sim. GBank gil586396 sp P38150 YB9Z_YEAST - HYPOTHETICAL 92.9 KD PROTEIN IN SSH1- APE3 INTERGENIC REGION		UNCLASSIFIED	1044
8314	30486802 (2801, 2802)	Novel Protein sim. GBank gil586398 sp P38210 YBA6_YEAST - HYPOTHETICAL 15.9 KD PROTEIN IN SAL1- PDR3 INTERGENIC REGION		UNCLASSIFIED	1038
8315	80026617 (6859, 6860)	Novel Protein sim. GBank gil586406 sp P38207 YBB9_YEAST - HYPOTHETICAL 59.4 KD PROTEIN IN RFT1- PEP1 INTERGENIC REGION		UNCLASSIFIED	1006, 1008, 1017, 1022, 1039

8316	29444896 (19095, 19096)	Novel Protein sim. GBank gil586406 sp P38207 YBB9_YEAST - HYPOTHETICAL 59.4 KD PROTEIN IN RFT1- PEP1 INTERGENIC REGION		UNCLASSIFIED	1022
8317	80223223 (18237, 18238)	Novel Protein sim. GBank gil586407 sp P38206 RFT1_YEAST - NUCLEAR DIVISION RFT1 PROTEIN		UNCLASSIFIED	1001, 1010, 1022, 1038
8318	8359887 (17655, 17656)	Novel Protein sim. GBank gil586426 sp P34216 YBE7_YEAST - HYPOTHETICAL 150.8 KD PROTEIN IN SEC17- QCR1 INTERGENIC REGION		UNCLASSIFIED	1024
8319	78739910 (627, 628)	Novel Protein sim. GBank gil586438 sp P38185 YBH1_YEAST - HYPOTHETICAL 11.5 KD PROTEIN IN RSP8A- AST1 INTERGENIC REGION		UNCLASSIFIED	1003, 1008, 1022, 1026
8320	79169341 (12817, 12818)	Novel Protein sim. GBank gil586455 sp P38170 YBJ7_YEAST - HYPOTHETICAL 83.0 KD PROTEIN IN ATP1- ROX3 INTERGENIC REGION		UNCLASSIFIED	1003, 1008, 1009, 1017, 1022, 1023, 1024, 1026, 1044
8321	20696454 (1859, 1860)	Novel Protein sim. GBank gil586458 sp P38167 YBK1_YEAST - HYPOTHETICAL 119.3 KD PROTEIN IN SFT2- ATP1 INTERGENIC REGION		UNCLASSIFIED	1022
8322	27213128 (8733, 8734)	Novel Protein sim. GBank gil586461 sp P38164 YBK4_YEAST - HYPOTHETICAL 104.7 KD PROTEIN IN PKC1- RTG3 INTERGENIC REGION		UNCLASSIFIED	1009
8323	29347099 (11371, 11372)	Novel Protein sim. GBank gil586461 sp P38164 YBK4_YEAST - HYPOTHETICAL 104.7 KD PROTEIN IN PKC1- RTG3 INTERGENIC REGION		UNCLASSIFIED	1009
8324	80217106 (3117, 3118)	Novel Protein sim. GBank gil586468 sp P38213 YBL7_YEAST - HYPOTHETICAL 82.4 KD PROTEIN IN COQ1- FLR1 INTERGENIC REGION		UNCLASSIFIED	1004, 1008, 1009, 1017, 1022, 1024, 1034, 1039

8325	27370166 (13451, 13452)	Novel Protein sim. GBank gi 586468 sp P38213 YBL7_YEAST - HYPOTHETICAL 82.4 KD PROTEIN IN COQ1- FLR1 INTERGENIC REGION		UNCLASSIFIED	1017
8326	80240850 (6529, 6530)	Novel Protein sim. GBank gi 586480 sp P38222 YBO0_YEAST - HYPOTHETICAL 62.6 KD PROTEIN IN CDS1- RPL2 INTERGENIC REGION		UNCLASSIFIED	1004, 1006, 1016, 1022, 1024, 1026, 1034, 1038
8327	17707791 (8747, 8748)	Novel Protein sim. GBank gi 586500 sp P38238 YBR1_YEAST - HYPOTHETICAL 34.7 KD PROTEIN IN ORC2- TIP1 INTERGENIC REGION		UNCLASSIFIED	1044
8328	80216068 (16415, 16416)	Novel Protein sim. GBank gi 586504 sp P38241 YBR5_YEAST - HYPOTHETICAL 40.9 KD PROTEIN IN ORC2- TIP1 INTERGENIC REGION		UNCLASSIFIED	1010, 1024, 1038, 1039
8329	94991988 (15923, 15924)	Novel Protein sim. GBank gi 586524 sp P38257 YBU8_YEAST - HYPOTHETICAL 52.8 KD PROTEIN IN VPS15- YMC2 INTERGENIC REGION		UNCLASSIFIED	1022
8330	27964479 (6285, 6286)	Novel Protein sim. GBank gi 586528 sp P38261 YBV2_YEAST - HYPOTHETICAL 85.5 KD PROTEIN IN VPS15- YMC2 INTERGENIC REGION		UNCLASSIFIED	1034
8331	80053850 (6833, 6834)	Novel Protein sim. GBank gi 586528 sp P38261 YBV2_YEAST - HYPOTHETICAL 85.5 KD PROTEIN IN VPS15- YMC2 INTERGENIC REGION		UNCLASSIFIED	1034
8332	78726368 (395, 396)	Novel Protein sim. GBank gi 586547 sp P38277 YBY8_YEAST - HYPOTHETICAL 60.5 KD PROTEIN IN ESR1- IRA1 INTERGENIC REGION		UNCLASSIFIED	1026
8333	79326867 (10575, 10576)	Novel Protein sim. GBank gi 586557 sp P37263 YC16_YEAST - HYPOTHETICAL 17.7 KD PROTEIN IN ABP1 5'REGION		UNCLASSIFIED	1003, 1004, 1022

8334	79568475 (5343, 5344)	Novel Protein sim. GBank gil586733 sp P37689 PMGL_ECOLI - PUTATIVE 2,3-BISPHOSPHOGLYCERATE-INDEPENDENT PHOSPHOGLYCERATE MUTASE (PHOSPHOGLYCEROMUTASE) (BPG- INDEPENDENT PGAM)		UNCLASSIFIED	1017
8335	16340393 (20383, 20384)	Novel Protein sim. GBank gil586848 sp P37518 YYAF_BACSU - HYPOTHETICAL 40.1 KD GTP-BINDING PROTEIN IN RPSF-SPO0J INTERGENIC REGION		UNCLASSIFIED	1022
8336	9667068 (1049, 1050)	Novel Protein sim. GBank gil586877 sp P37547 YABF_BACSU - HYPOTHETICAL 20.7 KD PROTEIN IN METS- KSGA INTERGENIC REGION		UNCLASSIFIED	1039
8337	15001498 (17119, 17120)	Novel Protein sim. GBank gil604483 (U18421) - Orf181 [Bacillus subtilis]		UNCLASSIFIED	1001
8338	78784791 (2453, 2454)	Novel Protein sim. GBank gil606451 emb CAA873671 - (Z47071) Ylm1p [Saccharomyces cerevisiae]		UNCLASSIFIED	1008, 1022
8339	20607081 (13613, 13614)	Novel Protein sim. GBank gil619595 (U17918) - unknown [Saccharomyces cerevisiae]		UNCLASSIFIED	1024
8340	94992079 (22303, 22304)	Novel Protein sim. GBank gil619595 (U17918) - unknown [Saccharomyces cerevisiae]		UNCLASSIFIED	1008, 1022, 1024, 1034
8341	79557452 (20113, 20114)	Novel Protein sim. GBank gil626442 pir S48545 - hypothetical protein YLR192c - yeast (Saccharomyces cerevisiae)		UNCLASSIFIED	1038
8342	78678607 (7905, 7906)	Novel Protein sim. GBank gil626443 pir S48546 - hypothetical protein YLR193c - yeast (Saccharomyces cerevisiae)		UNCLASSIFIED	1017
8343	56314582 (9849, 9850)	Novel Protein sim. GBank gil626444 pir S48547 - probable membrane protein YLR194c - yeast (Saccharomyces cerevisiae)		UNCLASSIFIED	1017, 1023, 1039, 1044
8344	27846851 (14553, 14554)	Novel Protein sim. GBank gil626659 pir S48981 - hypothetical protein YHR137w - yeast (Saccharomyces cerevisiae)		UNCLASSIFIED	1022

8345	27924944 (9401, 9402)	Novel Protein sim. GBank gij628869 pir S45894 - regulatory protein CSG2 precursor - yeast (Saccharomyces cerevisiae)	UNCLASSIFIED	1006
8346	27966665 (7895, 7896)	Novel Protein sim. GBank gij627071 pir A54523 - histidine-rich protein - Plasmodium lophurae (fragment)	UNCLASSIFIED	1022
8347	11669193 (7963, 7964)	Novel Protein sim. GBank gij627071 pir A54523 - histidine-rich protein - Plasmodium lophurae (fragment)	UNCLASSIFIED	1022
8348	87890190 (4259, 4260)	Novel Protein sim. GBank gij629282 pir S48578 - hypothetical protein - Mycoplasma capricolum (SGC3) (fragment)	UNCLASSIFIED	1008, 1015
8349	78674668 (18789, 18790)	Novel Protein sim. GBank gij642292 emb CAA87821 - (Z47815) unknown (Saccharomyces cerevisiae)	UNCLASSIFIED	1022
8350	20721189 (15105, 15106)	Novel Protein sim. GBank gij709805 gb AAC95474.1 - (U21300) transposase [Corynebacterium striatum]	UNCLASSIFIED	1006
8351	78780143 (20417, 20418)	Novel Protein sim. GBank gij710602 (L40588) - dynamin-related protein [Saccharomyces cerevisiae]	UNCLASSIFIED	1026
8352	30407707 (16389, 16390)	Novel Protein sim. GBank gij728659 emb CAA88527 - (Z48613) unknown (Saccharomyces cerevisiae)	UNCLASSIFIED	1008
8353	10237218 (1437, 1438)	Novel Protein sim. GBank gij728831 sp P39188 ALU1_HUMAN - IIII ALU SUBFAMILY J WARNING ENTRY IIII	UNCLASSIFIED	1025
8354	79245002 (4419, 4420)	Novel Protein sim. GBank gij728831 sp P39188 ALU1_HUMAN - IIII ALU SUBFAMILY J WARNING ENTRY IIII	UNCLASSIFIED	1008
8355	65883966 (13997, 13998)	Novel Protein sim. GBank gij728832 sp P39189 ALU2_HUMAN - IIII ALU SUBFAMILY SB WARNING ENTRY IIII	UNCLASSIFIED	1038
8356	78405952 (14535, 14536)	Novel Protein sim. GBank gij728834 sp P39191 ALU4_HUMAN - IIII ALU SUBFAMILY SB2 WARNING ENTRY IIII	UNCLASSIFIED	1026

8357	77527264 (2803, 2804)	Novel Protein sim. GBank gil728835 sp P39192 ALU5_HUMAN - III ALU SUBFAMILY SC WARNING ENTRY III		UNCLASSIFIED	1003
8358	86670623 (2917, 2918)	Novel Protein sim. GBank gil728835 sp P39192 ALU5_HUMAN - III ALU SUBFAMILY SC WARNING ENTRY III		UNCLASSIFIED	1049
8359	78512478 (17875, 17876)	Novel Protein sim. GBank gil728836 sp P39193 ALU6_HUMAN - III ALU SUBFAMILY SP WARNING ENTRY III		UNCLASSIFIED	1026
8360	15015938 (7163, 7164)	Novel Protein sim. GBank gil728838 sp P39195 ALU8_HUMAN - III ALU SUBFAMILY SX WARNING ENTRY III		UNCLASSIFIED	1054
8361	15021270 (18509, 18510)	Novel Protein sim. GBank gil728850 sp P08640 AMYH_YEAST - GLUCOAMYLASE S1/S2 PRECURSOR (GLUCAN 1,4-ALPHA-GLUCOSIDASE) (1,4- ALPHA-D-GLUCAN GLUCOHYDROLASE)		UNCLASSIFIED	1024
8362	85800306 (18913, 18914)	Novel Protein sim. GBank gil728850 sp P08640 AMYH_YEAST - GLUCOAMYLASE S1/S2 PRECURSOR (GLUCAN 1,4-ALPHA-GLUCOSIDASE) (1,4- ALPHA-D-GLUCAN GLUCOHYDROLASE)		UNCLASSIFIED	1025, 1034
8363	80412520 (11743, 11744)	Novel Protein sim. GBank gil728867 sp P40602 APG_ARATH - ANTER- SPECIFIC PROLINE-RICH PROTEIN APG - PRECURSOR		UNCLASSIFIED	1029
8364	80258491 (18067, 18068)	Novel Protein sim. GBank gil728867 sp P40602 APG_ARATH - ANTER- SPECIFIC PROLINE-RICH PROTEIN APG PRECURSOR		UNCLASSIFIED	1016, 1023, 1024
8365	27362122 (10033, 10034)	Novel Protein sim. GBank gil729138 sp P40987 CIN1_YEAST - CIN1 PROTEIN		UNCLASSIFIED	1026

8366	36995077 (6571, 6572)	Novel Protein sim. GBank gil729245 sp P40130 CYAA_ERWCH - ADENYLATE CYCLASE (ATP PYROPHOSPHATE-LYASE) (ADENYLAL CYCLASE)		UNCLASSIFIED	1016
8367	79247081 (5747, 5748)	Novel Protein sim. GBank gil729246 sp P40134 CYAA_HAEIN - ADENYLATE CYCLASE (ATP PYROPHOSPHATE-LYASE) (ADENYLAL CYCLASE)		UNCLASSIFIED	1008, 1022, 1026, 1039
8368	28491272 (7887, 7888)	Novel Protein sim. GBank gil729432 sp P22140 EPT1_YEAST - ETHANOLAMINEPHOSPHOTRANSFERASE (ETHPT)		UNCLASSIFIED	1008
8369	29264465 (6133, 6134)	Novel Protein sim. GBank gil730254 sp P40960 PA11_YEAST - PAC11 PROTEIN		UNCLASSIFIED	1022
8370	9051023 (18005, 18006)	Novel Protein sim. GBank gil730732 sp P32844 SEC6_YEAST - EXOCYST COMPLEX COMPONENT SEC6		UNCLASSIFIED	1006, 1010
8371	28477483 (18143, 18144)	Novel Protein sim. GBank gil730779 sp P38904 SP41_YEAST - SPP41 PROTEIN		UNCLASSIFIED	1008
8372	80212231 (13095, 13096)	Novel Protein sim. GBank gil731024 sp P38891 BCA1_YEAST - BRANCHED-CHAIN AMINO ACID AMINOTRANSFERASE, MITOCHONDRIAL PRECURSOR (BCAT) (TWT1 PROTEIN)		UNCLASSIFIED	1004, 1006, 1022
8373	5706051 (7973, 7974)	Novel Protein sim. GBank gil731317 sp P39559 YAM3_YEAST - HYPOTHETICAL 11.1 KD PROTEIN IN FLO1-PHO11 INTERGENIC REGION		UNCLASSIFIED	1058

8374	80217728 (17731, 17732)	Novel Protein sim. GBank gil731362 sp P38957 YD78_YEAST - HYPOTHETICAL 26.1 KD PROTEIN IN SED1- PDC2 INTERGENIC REGION		UNCLASSIFIED	1003, 1004, 1006, 1009, 1010, 1017, 1022, 1024, 1026, 1027, 1034, 1054
8375	8487223 (9751, 9752)	Novel Protein sim. GBank gil731388 sp P40103 YE18_YEAST - HYPOTHETICAL 27.3 KD PROTEIN IN ISC10 3'REGION		UNCLASSIFIED	1022
8376	21395476 (8431, 8432)	Novel Protein sim. GBank gil731397 sp P40001 YEA8_YEAST - HYPOTHETICAL 14.0 KD PROTEIN IN GCN4- WBP1 INTERGENIC REGION		UNCLASSIFIED	1006
8377	11611546 (20269, 20270)	Novel Protein sim. GBank gil731402 sp P39998 YEB5_YEAST - HYPOTHETICAL 61.3 KD PROTEIN IN PMP2- VAC8 INTERGENIC REGION		UNCLASSIFIED	1024
8378	71795603 (2849, 2850)	Novel Protein sim. GBank gil731408 sp P39993 YEC2_YEAST - HYPOTHETICAL 165.7 KD PROTEIN IN RIP1- URA3 INTERGENIC REGION		UNCLASSIFIED	1009
8379	78677913 (7777, 7778)	Novel Protein sim. GBank gil731435 sp P40009 YEJ5_YEAST - HYPOTHETICAL 71.9 KD PROTEIN IN PMI40- PAC2 INTERGENIC REGION		UNCLASSIFIED	1009
8380	85544655 (10757, 10758)	Novel Protein sim. GBank gil731435 sp P40009 YEJ5_YEAST - HYPOTHETICAL 71.9 KD PROTEIN IN PMI40- PAC2 INTERGENIC REGION		UNCLASSIFIED	1003, 1026, 1054
8381	30421541 (8517, 8518)	Novel Protein sim. GBank gil731445 sp P40017 YEL4_YEAST - HYPOTHETICAL 103.3 KD PROTEIN IN PRO3- GCD11 INTERGENIC REGION		UNCLASSIFIED	1026

8382	94239725 (4999, 5000)	Novel Protein sim. GBank gii731512 sp P40079 YEV7_YEAST - HYPOTHETICAL 40.8 KD PROTEIN IN RSP5- PAK1 INTERGENIC REGION		UNCLASSIFIED	1003
8383	80508435 (1165, 1166)	Novel Protein sim. GBank gii731515 sp P40081 YEW4_YEAST - HYPOTHETICAL 20.4 KD PROTEIN IN GLC7- GDI1 INTERGENIC REGION		UNCLASSIFIED	1004, 1006, 1012, 1017, 1044
8384	5705950 (3845, 3646)	Novel Protein sim. GBank gii731524 sp P40090 YEX7_YEAST - HYPOTHETICAL 72.1 KD PROTEIN IN FTR1- SPT15 INTERGENIC REGION		UNCLASSIFIED	1058
8385	27977136 (12789, 12790)	Novel Protein sim. GBank gii731528 sp P40093 YEV6_YEAST - HYPOTHETICAL 38.2 KD PROTEIN IN BEM2- NCB1 INTERGENIC REGION		UNCLASSIFIED	1022
8386	20709658 (14241, 14242)	Novel Protein sim. GBank gii731528 sp P40093 YEV6_YEAST - HYPOTHETICAL 38.2 KD PROTEIN IN BEM2- NCB1 INTERGENIC REGION		UNCLASSIFIED	1022
8387	80078984 (6731, 6732)	Novel Protein sim. GBank gii731576 sp P38888 YH04_YEAST - HYPOTHETICAL 91.2 KD PROTEIN IN RPS7A- SCH9 INTERGENIC REGION		UNCLASSIFIED	1001, 1008, 1022
8388	29012424 (1335, 1336)	Novel Protein sim. GBank gii731607 sp P38739 YHC8_YEAST - HYPOTHETICAL 63.8 KD PROTEIN IN GUT1- RIM1 INTERGENIC REGION PRECURSOR		UNCLASSIFIED	1017
8389	79415380 (20911, 20912)	Novel Protein sim. GBank gii731635 sp P38705 YHH1_YEAST - PUTATIVE SERYL-TRNA SYNTHETASE YHR011W (SERINE-TRNA LIGASE) (SERRS)		UNCLASSIFIED	1022
8390	30455534 (10763, 10764)	Novel Protein sim. GBank gii731676 sp P38796 YHN5_YEAST - HYPOTHETICAL 44.9 KD PROTEIN IN ERG7- NMD2 INTERGENIC REGION		UNCLASSIFIED	1008, 1022, 1026, 1039

8391	30671154 (11473, 11474)	Novel Protein sim. GBank gil731676 sp P38796 YHN5_YEAST - HYPOTHETICAL 44.9 KD PROTEIN IN ERG7- NMD2 INTERGENIC REGION		UNCLASSIFIED	1008
8392	29447086 (9079, 9080)	Novel Protein sim. GBank gil731679 sp P38800 YHO0_YEAST - HYPOTHETICAL 149.7 KD PROTEIN IN IRE1- KSP1 INTERGENIC REGION		UNCLASSIFIED	1026
8393	25243762 (22839, 22840)	Novel Protein sim. GBank gil731679 sp P38800 YHO0_YEAST - HYPOTHETICAL 149.7 KD PROTEIN IN IRE1- KSP1 INTERGENIC REGION		UNCLASSIFIED	1013
8394	28310311 (2481, 2482)	Novel Protein sim. GBank gil731707 sp P38827 SET1_YEAST - SET1 PROTEIN		UNCLASSIFIED	1003
8395	57104763 (14275, 14276)	Novel Protein sim. GBank gil731726 sp P32900 YHU9_YEAST - HYPOTHETICAL 81.8 KD PROTEIN IN MRPL6- SPO12 INTERGENIC REGION		UNCLASSIFIED	1024
8396	28780684 (10301, 10302)	Novel Protein sim. GBank gil731729 sp P38850 YHV4_YEAST - HYPOTHETICAL 123.0 KD PROTEIN IN SPO16- REC104 INTERGENIC REGION		UNCLASSIFIED	1022
8397	82124927 (14327, 14328)	Novel Protein sim. GBank gil731729 sp P38850 YHV4_YEAST - HYPOTHETICAL 123.0 KD PROTEIN IN SPO16- REC104 INTERGENIC REGION		UNCLASSIFIED	1004, 1006, 1010, 1012, 1016, 1022, 1024, 1034, 1038, 1044
8398	85541496 (5329, 5330)	Novel Protein sim. GBank gil731743 sp P38863 YHX2_YEAST - HYPOTHETICAL 96.8 KD PROTEIN IN NMD3- ENO2 INTERGENIC REGION		UNCLASSIFIED	1038
8399	79831968 (8349, 8350)	Novel Protein sim. GBank gil731743 sp P38863 YHX2_YEAST - HYPOTHETICAL 96.8 KD PROTEIN IN NMD3- ENO2 INTERGENIC REGION		UNCLASSIFIED	1006, 1008, 1010, 1017, 1034

8400	82056504 (19157, 19158)	Novel Protein sim. GBank gil731744 sp P38864 YHX3_YEAST - HYPOTHETICAL 12.9 KD PROTEIN IN NMD3- ENO2 INTERGENIC REGION		UNCLASSIFIED	1004, 1006, 1008, 1010, 1012, 1016, 1022, 1023, 1024, 1025, 1034, 1038, 1039, 1049
8401	78912783 (21427, 21428)	Novel Protein sim. GBank gil731751 sp P38868 YHY0_YEAST - HYPOTHETICAL 18.1 KD PROTEIN IN OYE2- GND1 INTERGENIC REGION		UNCLASSIFIED	1009, 1023
8402	37803858 (10587, 10588)	Novel Protein sim. GBank gil731753 sp P38870 YHY2_YEAST - HYPOTHETICAL 90.1 KD PROTEIN IN OYE2- GND1 INTERGENIC REGION		UNCLASSIFIED	1058
8403	57531138 (5309, 5310)	Novel Protein sim. GBank gil731784 sp P40547 YIB7_YEAST - HYPOTHETICAL 105.5 KD PROTEIN IN RPL5A- BAR1 INTERGENIC REGION		UNCLASSIFIED	1001, 1004, 1006, 1010, 1012, 1022, 1023, 1024, 1031, 1034
8404	79861578 (14117, 14118)	Novel Protein sim. GBank gil731812 sp P40524 YIF4_YEAST - HYPOTHETICAL 12.0 KD PROTEIN IN SNP1- GPP1 INTERGENIC REGION		UNCLASSIFIED	1017, 1025, 1027
8405	80233573 (17473, 17474)	Novel Protein sim. GBank gil731823 sp P40515 YIG6_YEAST - HYPOTHETICAL 17.7 KD PROTEIN IN RNR3- ARC15 INTERGENIC REGION		UNCLASSIFIED	1004, 1006, 1008, 1017, 1022, 1023, 1034
8406	28773835 (8341, 8342)	Novel Protein sim. GBank gil731836 sp P40504 KTR7_YEAST - PROBABLE MANNOSYLTRANSFERASE KTR7		UNCLASSIFIED	1023
8407	11395459 (12907, 12908)	Novel Protein sim. GBank gil731841 sp P40499 YIJ0_YEAST - HYPOTHETICAL 56.3 KD PROTEIN IN SGA1- KTR7 INTERGENIC REGION		UNCLASSIFIED	1006

8408	71799569 (15059, 15060)	Novel Protein sim. GBank gil731844 sp P40496 YIJ3_YEAST - HYPOTHETICAL 30.5 KD PROTEIN IN SGA1- KTR7 INTERGENIC REGION		UNCLASSIFIED	1034
8409	11614694 (18465, 18466)	Novel Protein sim. GBank gil731847 sp P40492 YIJ7_YEAST - HYPOTHETICAL 59.9 KD PROTEIN IN SGA1- KTR7 INTERGENIC REGION		UNCLASSIFIED	1034
8410	57293944 (18147, 18148)	Novel Protein sim. GBank gil731852 sp P40487 YIK3_YEAST - HYPOTHETICAL 48.3 KD PROTEIN IN MOB1- SGA1 INTERGENIC REGION		UNCLASSIFIED	1004, 1008, 1010, 1022, 1024
8411	78619106 (15633, 15634)	Novel Protein sim. GBank gil731871 sp P40468 YIM9_YEAST - HYPOTHETICAL 269.9 KD PROTEIN IN FKH1- MET18 INTERGENIC REGION		UNCLASSIFIED	1006, 1025
8412	85053334 (1137, 1138)	Novel Protein sim. GBank gil731916 sp P40572 YIT6_YEAST - HYPOTHETICAL 28.9 KD PROTEIN IN DBF8- MET28 INTERGENIC REGION		UNCLASSIFIED	1024
8413	80025402 (14749, 14750)	Novel Protein sim. GBank gil731938 sp P40364 YJG6_YEAST - HYPOTHETICAL 28.5 KD PROTEIN IN SMC3- MRPL8 INTERGENIC REGION		UNCLASSIFIED	1004
8414	80052979 (14721, 14722)	Novel Protein sim. GBank gil731956 sp P39541 YJT5_YEAST - HYPOTHETICAL 25.4 KD PROTEIN IN UBP12- CDC6 INTERGENIC REGION		UNCLASSIFIED	1017, 1024, 1054
8415	16700856 (13687, 13688)	Novel Protein sim. GBank gil731975 sp P40858 YJG6_YEAST - HYPOTHETICAL 25.4 KD PROTEIN IN SAP185- BCK1 INTERGENIC REGION		UNCLASSIFIED	1008, 1022
8416	78928384 (577, 578)	Novel Protein sim. GBank gil732094 sp P39386 YJIO_ECOLI - HYPOTHETICAL 44.7 KD PROTEIN IN IADA- MCRD INTERGENIC REGION (F410)		UNCLASSIFIED	1008

8417	80095666 (15429, 15430)	Novel Protein sim. GBank gil732099 sp P39391 YJIT_ECOLI - HYPOTHETICAL 58.6 KD PROTEIN IN IADA- MCRD INTERGENIC REGION	UNCLASSIFIED	1006, 1008, 1024, 1044
8418	10131805 (22871, 22872)	Novel Protein sim. GBank gil732099 sp P39391 YJIT_ECOLI - HYPOTHETICAL 58.6 KD PROTEIN IN IADA- MCRD INTERGENIC REGION	UNCLASSIFIED	1017
8419	80252185 (17285, 17286)	Novel Protein sim. GBank gil732111 sp P39399 YJIM_ECOLI - HYPOTHETICAL 31.4 KD PROTEIN IN TSR- MDOB INTERGENIC REGION (F288B)	UNCLASSIFIED	1004, 1039
8420	80244108 (8381, 8382)	Novel Protein sim. GBank gil732162 sp P40214 YM22_YEAST - HYPOTHETICAL 38.6 KD PROTEIN IN RPS16A- TIF34 INTERGENIC REGION	UNCLASSIFIED	1023, 1034, 1044, 1054
8421	78459704 (12195, 12196)	Novel Protein sim. GBank gil732185 sp P40156 YINV3_YEAST - HYPOTHETICAL 25.3 KD PROTEIN IN PEX17- MER1 INTERGENIC REGION	UNCLASSIFIED	1026
8422	17699503 (21589, 21590)	Novel Protein sim. GBank gil746496 (U23515) - weak similarity to adenyl cyclase-associated protein (CAP) and to P. chabaudi adami major merozoite surface antigen protein (PIR:A32555). Final exon overlaps gene predicted on other strand. [Caenorhabditis elegans]	UNCLASSIFIED	1003
8423	80079362 (8457, 8458)	Novel Protein sim. GBank gil76177 pir QQECFT - hypothetical 38.8K protein (ftsI 5' region) - Escherichia coli	UNCLASSIFIED	1034
8424	8908046 (8165, 8166)	Novel Protein sim. GBank gil76188 pir QQECP7 - hypothetical 7.6K protein (pbpA 5' region) - Escherichia coli	UNCLASSIFIED	1017
8425	25316189 (19767, 19768)	Novel Protein sim. GBank gil78586 pir C24833 - hypothetical protein 2 - Escherichia coli (fragment)	UNCLASSIFIED	1009

8426	78462250 (6063, 6064)	Novel Protein sim. GBank gil798882 emb CAA89100 - (Z49210) unknown [Saccharomyces cerevisiae]		UNCLASSIFIED	1008
8427	28971889 (14281, 14282)	Novel Protein sim. GBank gil805026 emb CAA89163 - (Z49219) unknown [Saccharomyces cerevisiae]		UNCLASSIFIED	1022
8428	65702316 (9507, 9508)	Novel Protein sim. GBank gil81286 pir S22697 - extensin - Volvox carteri (fragment)		UNCLASSIFIED	1038
8429	80199647 (907, 908)	Novel Protein sim. GBank gil81869 pir C29356 - hydroxyproline-rich glycoprotein (clone Hyp2.13) - kidney bean (fragment)		UNCLASSIFIED	1034
8430	78478086 (20397, 20398)	Novel Protein sim. GBank gil81869 pir C29356 - hydroxyproline-rich glycoprotein (clone Hyp2.13) - kidney bean (fragment)		UNCLASSIFIED	1003, 1039
8431	88095247 (19101, 19102)	Novel Protein sim. GBank gil83715 pir JH0262 - QutR protein - Emericella nidulans		UNCLASSIFIED	1034
8432	10143820 (621, 622)	Novel Protein sim. GBank gil854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1054
8433	10168378 (699, 700)	Novel Protein sim. GBank gil854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1025
8434	7520561 (1675, 1676)	Novel Protein sim. GBank gil854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1058
8435	16314987 (3203, 3204)	Novel Protein sim. GBank gil854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1054
8436	57294076 (4829, 4830)	Novel Protein sim. GBank gil854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1034
8437	35140774 (5295, 5296)	Novel Protein sim. GBank gil854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1050
8438	19426288 (5451, 5452)	Novel Protein sim. GBank gil854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1054

8439	80391876 (6391, 6392)	Novel Protein sim. GBank gi 854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1023, 1029, 1034, 1038, 1044
8440	15005436 (6559, 6560)	Novel Protein sim. GBank gi 854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1027
8441	39731954 (7735, 7736)	Novel Protein sim. GBank gi 854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1026
8442	78053184 (9363, 9364)	Novel Protein sim. GBank gi 854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1026
8443	11399483 (11915, 11916)	Novel Protein sim. GBank gi 854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1024
8444	80245970 (12473, 12474)	Novel Protein sim. GBank gi 854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1053
8445	80441138 (13101, 13102)	Novel Protein sim. GBank gi 854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1038, 1039
8446	80058938 (16345, 16346)	Novel Protein sim. GBank gi 854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1010
8447	80226511 (16939, 16940)	Novel Protein sim. GBank gi 854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1034
8448	87463001 (16995, 16996)	Novel Protein sim. GBank gi 854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1015
8449	80433688 (17269, 17270)	Novel Protein sim. GBank gi 854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1034, 1038, 1039
8450	34882457 (17623, 17624)	Novel Protein sim. GBank gi 854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1000

8451	20628223 (19323, 19324)	Novel Protein sim. GBank gi 854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1034
8452	79240757 (19455, 19456)	Novel Protein sim. GBank gi 854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1008
8453	79420403 (22213, 22214)	Novel Protein sim. GBank gi 854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1034
8454	80023287 (22431, 22432)	Novel Protein sim. GBank gi 854065 emb CAA58337 - (X83413) U88 [Human herpesvirus 6]		UNCLASSIFIED	1010, 1034
8455	37800896 (8987, 8988)	Novel Protein sim. GBank gi 860970 emb CAA60698 - (X87248) HP8 peptide [Homo sapiens]		UNCLASSIFIED	1012
8456	80306282 (7295, 7296)	Novel Protein sim. GBank gi 868023 (M14641) - ORF downstream of manA [Escherichia coli]		UNCLASSIFIED	1017, 1029
8457	21425808 (967, 968)	Novel Protein sim. GBank gi 882648 (U29579) - ORF_f305 [Escherichia coli]		UNCLASSIFIED	1022
8458	29013247 (15587, 15588)	Novel Protein sim. GBank gi 887600 emb CAA90190 - (Z49939) unknown [Saccharomyces cerevisiae]		UNCLASSIFIED	1039
8459	20446447 (12205, 12206)	Novel Protein sim. GBank gi 887831 (U28375) - ORF_0956 [Escherichia coli]		UNCLASSIFIED	1024
8460	20128919 (2081, 2082)	Novel Protein sim. GBank gi 905361 (U22103) - gag-protease polyprotein [Glycine max]		UNCLASSIFIED	1034
8461	79970410 (7843, 7844)	Novel Protein sim. GBank gi 92776 pir B29545 - submandibular gland basic polypeptide - rat		UNCLASSIFIED	1024
8462	21416803 (20445, 20446)	Novel Protein sim. GBank gi 93505 pir B34768 - ORF5 protein - Orf virus (strain NZ2)		UNCLASSIFIED	1034
8463	66437111 (22329, 22330)	Novel Protein sim. GBank gi 95799 pir S18536 - envelope protein - Escherichia coli		UNCLASSIFIED	1024
8464	66396557 (20973, 20974)	Novel Protein sim. GBank gi 97480 pir S19739 - integral membrane protein - Rhodobacter capsulatus		UNCLASSIFIED	1008
8465	29203958 (12991, 12992)	Novel Protein sim. GBank gi 984814 (U27196) - zinc finger protein [Gallus gallus]		UNCLASSIFIED	1017

8466	19485564 (9043, 9044)	Novel Protein sim. GBank gij987652[dbj BAA10924] - (D64044) hypothylcal protein in UNG 3' region [Escherichia coli]		UNCLASSIFIED	1016
8467	79618357 (6783, 6784)	Novel Protein sim. GBank gij98837[pir A41326 - enantiomer-selective amidase - Rhodococcus sp		UNCLASSIFIED	1039
8468	78464431 (19939, 19940)	Novel Protein sim. GBank gij2821973[dbj BAA24537] - (AB010100) aquaporin 7 [Mus musculus]	Contains protein domain (PF00230) - Major intrinsic protein	water_channel	1003
8469	71787672 (20291, 20292)	Novel Protein sim. GBank gij3264596 (AF057183) - putative tonoplast aquaporin [Zea mays]	Contains protein domain (PF00230) - Major intrinsic protein	water_channel	1053
8470	19888144 (7, 8)			UNCLASSIFIED	1034
8471	50100675 (9, 10)			UNCLASSIFIED	1036
8472	17905278 (29, 30)			UNCLASSIFIED	1024
8473	78790953 (43, 44)			UNCLASSIFIED	1009, 1010, 1017, 1039
8474	8491147 (55, 56)			UNCLASSIFIED	1034
8475	17087218 (57, 58)			UNCLASSIFIED	1023
8476	16289836 (71, 72)			UNCLASSIFIED	1054
8477	28987802 (97, 98)			UNCLASSIFIED	1022
8478	38438928 (105, 106)			UNCLASSIFIED	1034
8479	79111647 (107, 108)			UNCLASSIFIED	1044
8480	46866607 (119, 120)			UNCLASSIFIED	1037
8481	38069815 (123, 124)			UNCLASSIFIED	1019
8482	27965974 (143, 144)			UNCLASSIFIED	1006

8483	20721254 (165, 166)			UNCLASSIFIED	1006
8484	80019493 (183, 184)			UNCLASSIFIED	1022, 1034
8485	86468733 (187, 188)			UNCLASSIFIED	1029
8486	30243976 (199, 200)			UNCLASSIFIED	1026
8487	27932090 (209, 210)			UNCLASSIFIED	1022
8488	10200065 (211, 212)			UNCLASSIFIED	1025
8489	11424810 (219, 220)			UNCLASSIFIED	1024
8490	65888031 (253, 254)			UNCLASSIFIED	1054
8491	80537645 (257, 258)			UNCLASSIFIED	1024, 1058
8492	23290881 (259, 260)			UNCLASSIFIED	1012, 1017, 1023
8493	85805467 (279, 280)			UNCLASSIFIED	1054
8494	65451689 (281, 282)			UNCLASSIFIED	1016
8495	79852910 (283, 284)			UNCLASSIFIED	1003, 1044, 1058
8496	11617297 (289, 290)			UNCLASSIFIED	1024
8497	79774533 (293, 294)			UNCLASSIFIED	1003, 1023, 1025, 1034, 1044, 1054
8498	10119777 (301, 302)			UNCLASSIFIED	1044
8499	32290705 (311, 312)			UNCLASSIFIED	1000
8500	66365477 (313, 314)			UNCLASSIFIED	1054

8501	87113696 (317, 318)			UNCLASSIFIED	1007, 1018
8502	19649670 (319, 320)			UNCLASSIFIED	1023
8503	29347217 (333, 334)			UNCLASSIFIED	1017
8504	50673835 (335, 336)			UNCLASSIFIED	1036, 1037
8505	82417404 (349, 350)			UNCLASSIFIED	1001, 1010, 1022
8506	5943483 (351, 352)			UNCLASSIFIED	1058
8507	94323836 (365, 366)			UNCLASSIFIED	1054
8508	27988090 (369, 370)			UNCLASSIFIED	1054
8509	16298783 (371, 372)			UNCLASSIFIED	1054
8510	78171074 (377, 378)			UNCLASSIFIED	1050
8511	38922194 (379, 380)			UNCLASSIFIED	1008
8512	91237148 (385, 386)			UNCLASSIFIED	1045
8513	11776386 (387, 388)			UNCLASSIFIED	1006
8514	37057869 (405, 406)			UNCLASSIFIED	1054
8515	47656029 (415, 416)			UNCLASSIFIED	1029
8516	20709254 (419, 420)			UNCLASSIFIED	1022
8517	54697416 (431, 432)			UNCLASSIFIED	1022
8518	30788671 (435, 436)			UNCLASSIFIED	1003

8519	79172297 (439, 440)			UNCLASSIFIED	1024
8520	87461475 (457, 458)			UNCLASSIFIED	1025
8521	56511989 (463, 464)			UNCLASSIFIED	1046
8522	80067736 (465, 466)			UNCLASSIFIED	1017, 1022
8523	35351646 (473, 474)			UNCLASSIFIED	1016
8524	10116015 (481, 482)			UNCLASSIFIED	1023
8525	37057574 (491, 492)			UNCLASSIFIED	1054
8526	79580738 (497, 498)			UNCLASSIFIED	1024, 1025
8527	81895393 (501, 502)			UNCLASSIFIED	1008, 1017, 1034, 1038, 1054
8528	46889925 (503, 504)			UNCLASSIFIED	1037
8529	79969257 (509, 510)			UNCLASSIFIED	1010
8530	79635783 (517, 518)			UNCLASSIFIED	1016, 1038
8531	65455454 (519, 520)			UNCLASSIFIED	1016
8532	20273021 (533, 534)			UNCLASSIFIED	1022
8533	8761610 (535, 536)			UNCLASSIFIED	1004
8534	78742780 (547, 548)			UNCLASSIFIED	1044
8535	29687766 (551, 552)			UNCLASSIFIED	1024
8536	86673999 (555, 556)			UNCLASSIFIED	1007

8537	57056453 (563, 564)			UNCLASSIFIED	1054
8538	66557903 (565, 566)			UNCLASSIFIED	1024, 1029
8539	20287934 (571, 572)			UNCLASSIFIED	1022
8540	94851589 (583, 584)			UNCLASSIFIED	1010, 1024, 1034
8541	33267436 (599, 600)			UNCLASSIFIED	1001
8542	55410094 (609, 610)			UNCLASSIFIED	1021
8543	66726855 (619, 620)			UNCLASSIFIED	1009
8544	65709003 (629, 630)			UNCLASSIFIED	1023
8545	79567455 (631, 632)			UNCLASSIFIED	1008
8546	33182618 (637, 638)			UNCLASSIFIED	1026
8547	70628031 (639, 640)			UNCLASSIFIED	1036
8548	11102356 (643, 644)			UNCLASSIFIED	1053
8549	23289089 (645, 646)			UNCLASSIFIED	1003
8550	29687952 (653, 654)			UNCLASSIFIED	1024
8551	55663214 (655, 656)			UNCLASSIFIED	1021
8552	70623103 (663, 664)			UNCLASSIFIED	1019, 1036
8553	27966627 (675, 676)			UNCLASSIFIED	1053
8554	36621832 (681, 682)			UNCLASSIFIED	1034

8555	65857003 (685, 686)			UNCLASSIFIED	1038
8556	11011418 (689, 690)			UNCLASSIFIED	1024
8557	65492939 (695, 696)			UNCLASSIFIED	1008
8558	87132885 (697, 698)			UNCLASSIFIED	1025
8559	33187779 (709, 710)			UNCLASSIFIED	1026
8560	87451834 (711, 712)			UNCLASSIFIED	1025
8561	65578630 (725, 726)			UNCLASSIFIED	1016, 1046
8562	77713567 (737, 738)			UNCLASSIFIED	1054
8563	30279273 (741, 742)			UNCLASSIFIED	1050
8564	57118476 (743, 744)			UNCLASSIFIED	1054
8565	32275288 (761, 762)			UNCLASSIFIED	1029
8566	11600280 (791, 792)			UNCLASSIFIED	1006
8567	86668944 (797, 798)			UNCLASSIFIED	1007
8568	80085067 (799, 800)			UNCLASSIFIED	1001
8569	78679168 (811, 812)			UNCLASSIFIED	1009
8570	29361610 (821, 822)			UNCLASSIFIED	1009
8571	78455843 (837, 838)			UNCLASSIFIED	1008
8572	66468628 (857, 858)			UNCLASSIFIED	1019, 1024

8573	11696012 (859, 860)			UNCLASSIFIED	1034
8574	80052094 (865, 866)			UNCLASSIFIED	1010, 1024
8575	65649526 (889, 890)			UNCLASSIFIED	1038
8576	11093325 (897, 898)			UNCLASSIFIED	1022
8577	33765010 (899, 900)			UNCLASSIFIED	1026
8578	11804718 (901, 902)			UNCLASSIFIED	1038
8579	27966751 (913, 914)			UNCLASSIFIED	1022
8580	38896195 (921, 922)			UNCLASSIFIED	1024
8581	79250611 (923, 924)			UNCLASSIFIED	1024, 1026
8582	85539103 (925, 926)			UNCLASSIFIED	1054
8583	80243138 (931, 932)			UNCLASSIFIED	1022
8584	78444716 (933, 934)			UNCLASSIFIED	1026
8585	20292771 (935, 936)			UNCLASSIFIED	1034
8586	30167384 (941, 942)			UNCLASSIFIED	1030
8587	37811964 (943, 944)			UNCLASSIFIED	1058
8588	66367854 (957, 958)			UNCLASSIFIED	1004
8589	47652097 (959, 960)			UNCLASSIFIED	1029
8590	29232603 (963, 964)			UNCLASSIFIED	1017

8591	36509035 (969, 970)			UNCLASSIFIED	1026, 1029
8592	66049560 (987, 988)			UNCLASSIFIED	1007, 1024
8593	66049699 (989, 990)			UNCLASSIFIED	1054
8594	11799492 (1007, 1008)			UNCLASSIFIED	1038
8595	30787062 (1021, 1022)			UNCLASSIFIED	1026
8596	10310325 (1033, 1034)			UNCLASSIFIED	1038
8597	55178182 (1047, 1048)			UNCLASSIFIED	1019
8598	11386206 (1051, 1052)			UNCLASSIFIED	1016
8599	33202361 (1053, 1054)			UNCLASSIFIED	1026
8600	21427719 (1055, 1056)			UNCLASSIFIED	1022
8601	77663286 (1057, 1058)			UNCLASSIFIED	1038
8602	79371675 (1061, 1062)			UNCLASSIFIED	1009, 1010, 1022, 1024
8603	98065610 (1065, 1066)			UNCLASSIFIED	1010
8604	94147083 (1069, 1070)			UNCLASSIFIED	1003
8605	19402999 (1071, 1072)			UNCLASSIFIED	1025
8606	54860382 (1081, 1082)			UNCLASSIFIED	1019
8607	18469943 (1083, 1084)			UNCLASSIFIED	1054
8608	55005583 (1095, 1096)			UNCLASSIFIED	1019

8609	78755953 (1099, 1100)			UNCLASSIFIED	1022
8610	46863455 (1107, 1108)			UNCLASSIFIED	1050
8611	42724831 (1117, 1118)			UNCLASSIFIED	1000
8612	94846402 (1119, 1120)			UNCLASSIFIED	1044
8613	71091917 (1143, 1144)			UNCLASSIFIED	1053
8614	8485735 (1185, 1186)			UNCLASSIFIED	1004
8615	38916526 (1193, 1194)			UNCLASSIFIED	1026
8616	78361234 (1197, 1198)			UNCLASSIFIED	1008, 1026
8617	70947442 (1211, 1212)			UNCLASSIFIED	1029
8618	65876990 (1215, 1216)			UNCLASSIFIED	1038
8619	78369099 (1221, 1222)			UNCLASSIFIED	1039
8620	10357663 (1223, 1224)			UNCLASSIFIED	1039
8621	55181143 (1229, 1230)			UNCLASSIFIED	1019
8622	55482059 (1231, 1232)			UNCLASSIFIED	1026
8623	57311213 (1245, 1246)			UNCLASSIFIED	1054
8624	30271112 (1249, 1250)			UNCLASSIFIED	1008
8625	79399248 (1271, 1272)			UNCLASSIFIED	1024
8626	19894263 (1279, 1280)			UNCLASSIFIED	1022

8627	33184739 (1283, 1284)			UNCLASSIFIED	1026
8628	29224802 (1297, 1298)			UNCLASSIFIED	1030
8629	78500861 (1299, 1300)			UNCLASSIFIED	1054
8630	33202266 (1301, 1302)			UNCLASSIFIED	1026
8631	78082996 (1311, 1312)			UNCLASSIFIED	1029
8632	55773013 (1317, 1318)			UNCLASSIFIED	1021
8633	46743047 (1319, 1320)			UNCLASSIFIED	1034
8634	32612112 (1341, 1342)			UNCLASSIFIED	1029
8635	94656612 (1343, 1344)			UNCLASSIFIED	1004, 1024
8636	98422253 (1349, 1350)			UNCLASSIFIED	1003, 1020
8637	79390676 (1355, 1356)			UNCLASSIFIED	1058
8638	16292585 (1377, 1378)			UNCLASSIFIED	1054
8639	86885799 (1383, 1384)			UNCLASSIFIED	1029, 1054
8640	66364813 (1403, 1404)			UNCLASSIFIED	1004
8641	65445469 (1411, 1412)			UNCLASSIFIED	1038
8642	35907435 (1413, 1414)			UNCLASSIFIED	1029
8643	78374001 (1419, 1420)			UNCLASSIFIED	1003
8644	27371637 (1421, 1422)			UNCLASSIFIED	1022

8645	77822646 (1423, 1424)			UNCLASSIFIED	1022
8646	54530230 (1429, 1430)			UNCLASSIFIED	1041
8647	33203723 (1433, 1434)			UNCLASSIFIED	1026
8648	86671304 (1435, 1436)			UNCLASSIFIED	1007
8649	71761926 (1443, 1444)			UNCLASSIFIED	1000, 1008
8650	30666980 (1445, 1446)			UNCLASSIFIED	1008
8651	11413565 (1449, 1450)			UNCLASSIFIED	1017
8652	37802417 (1453, 1454)			UNCLASSIFIED	1058
8653	38903943 (1455, 1456)			UNCLASSIFIED	1026
8654	13051575 (1459, 1460)			UNCLASSIFIED	1027
8655	46852581 (1461, 1462)			UNCLASSIFIED	1037
8656	20457546 (1463, 1464)			UNCLASSIFIED	1010
8657	54527382 (1485, 1486)			UNCLASSIFIED	1041
8658	20436589 (1493, 1494)			UNCLASSIFIED	1010
8659	25258276 (1495, 1496)			UNCLASSIFIED	1039
8660	66556694 (1497, 1498)			UNCLASSIFIED	1008
8661	80251683 (1499, 1500)			UNCLASSIFIED	1010
8662	65498691 (1505, 1506)			UNCLASSIFIED	1016

8663	7785453 (1517, 1518)			UNCLASSIFIED	1021
8664	77932495 (1523, 1524)			UNCLASSIFIED	1036
8665	79765487 (1529, 1530)			UNCLASSIFIED	1022, 1025
8666	11798274 (1531, 1532)			UNCLASSIFIED	1038
8667	91230900 (1533, 1534)			UNCLASSIFIED	1010
8668	94238509 (1539, 1540)			UNCLASSIFIED	1010
8669	78269661 (1555, 1556)			UNCLASSIFIED	1006, 1054
8670	80424153 (1563, 1564)			UNCLASSIFIED	1022, 1025
8671	85803274 (1569, 1570)			UNCLASSIFIED	1054
8672	16839473 (1571, 1572)			UNCLASSIFIED	1023
8673	79772287 (1579, 1580)			UNCLASSIFIED	1003
8674	19543657 (1581, 1582)			UNCLASSIFIED	1001
8675	46866612 (1589, 1590)			UNCLASSIFIED	1037
8676	30265081 (1605, 1606)			UNCLASSIFIED	1050
8677	21145664 (1623, 1624)			UNCLASSIFIED	1058
8678	65868045 (1633, 1634)			UNCLASSIFIED	1016
8679	25335978 (1639, 1640)			UNCLASSIFIED	1003
8680	9894247 (1647, 1648)			UNCLASSIFIED	1044

8681	30522206 (1651, 1652)			UNCLASSIFIED	1016
8682	37798922 (1657, 1658)			UNCLASSIFIED	1012
8683	78406747 (1667, 1668)			UNCLASSIFIED	1024
8684	55177260 (1695, 1696)			UNCLASSIFIED	1019
8685	56026288 (1699, 1700)			UNCLASSIFIED	1009
8686	71215348 (1721, 1722)			UNCLASSIFIED	1029
8687	78077450 (1725, 1726)			UNCLASSIFIED	1029
8688	78902979 (1735, 1736)			UNCLASSIFIED	1003
8689	30797675 (1737, 1738)			UNCLASSIFIED	1026
8690	17658790 (1739, 1740)			UNCLASSIFIED	1017
8691	65888040 (1749, 1750)			UNCLASSIFIED	1054
8692	77880626 (1753, 1754)			UNCLASSIFIED	1019
8693	55006873 (1771, 1772)			UNCLASSIFIED	1019
8694	79955700 (1777, 1778)			UNCLASSIFIED	1024, 1025
8695	79880032 (1793, 1794)			UNCLASSIFIED	1017
8696	94852145 (1813, 1814)			UNCLASSIFIED	1010
8697	78891791 (1835, 1836)			UNCLASSIFIED	1003
8698	78772065 (1839, 1840)			UNCLASSIFIED	1008

8699	29000390 (1863, 1864)			UNCLASSIFIED	1030
8700	78524207 (1867, 1868)			UNCLASSIFIED	1003
8701	21428127 (1877, 1878)			UNCLASSIFIED	1022
8702	78747624 (1879, 1880)			UNCLASSIFIED	1026
8703	36222246 (1885, 1886)			UNCLASSIFIED	1034
8704	65465725 (1889, 1890)			UNCLASSIFIED	1054
8705	79372584 (1893, 1894)			UNCLASSIFIED	1000, 1024
8706	29679359 (1901, 1902)			UNCLASSIFIED	1038
8707	11286739 (1913, 1914)			UNCLASSIFIED	1006
8708	79603242 (1915, 1916)			UNCLASSIFIED	1003, 1044
8709	19517549 (1927, 1928)			UNCLASSIFIED	1044
8710	30650824 (1949, 1950)			UNCLASSIFIED	1008
8711	17927742 (1955, 1956)			UNCLASSIFIED	1038
8712	80421364 (1957, 1958)			UNCLASSIFIED	1025
8713	17087222 (1959, 1960)			UNCLASSIFIED	1023
8714	65675571 (2003, 2004)			UNCLASSIFIED	1046
8715	29013650 (2007, 2008)			UNCLASSIFIED	1030
8716	78782114 (2011, 2012)			UNCLASSIFIED	1009

8717	87456028 (2013, 2014)			UNCLASSIFIED	1025
8718	16554210 (2017, 2018)			UNCLASSIFIED	1022, 1023
8719	23284764 (2019, 2020)			UNCLASSIFIED	1039
8720	94317594 (2027, 2028)			UNCLASSIFIED	1044
8721	65648570 (2033, 2034)			UNCLASSIFIED	1054
8722	28992438 (2039, 2040)			UNCLASSIFIED	1030
8723	78923542 (2041, 2042)			UNCLASSIFIED	1054
8724	65707106 (2049, 2050)			UNCLASSIFIED	1016
8725	65983890 (2051, 2052)			UNCLASSIFIED	1004
8726	20589930 (2063, 2064)			UNCLASSIFIED	1053
8727	65992108 (2073, 2074)			UNCLASSIFIED	1016, 1044
8728	54994410 (2087, 2088)			UNCLASSIFIED	1029
8729	39530223 (2101, 2102)			UNCLASSIFIED	1022
8730	46780980 (2105, 2106)			UNCLASSIFIED	1036
8731	79836592 (2115, 2116)			UNCLASSIFIED	1008, 1044, 1053
8732	52559484 (2119, 2120)			UNCLASSIFIED	1058
8733	85805476 (2121, 2122)			UNCLASSIFIED	1054

8734	80083435 (2131, 2132)			UNCLASSIFIED	1001, 1003, 1008, 1009, 1022, 1023, 1026, 1044
8735	37017348 (2133, 2134)			UNCLASSIFIED	1025
8736	78528971 (2141, 2142)			UNCLASSIFIED	1026
8737	11100463 (2147, 2148)			UNCLASSIFIED	1022
8738	24141933 (2183, 2184)			UNCLASSIFIED	1003
8739	19631713 (2185, 2186)			UNCLASSIFIED	1004
8740	78280739 (2191, 2192)			UNCLASSIFIED	1000
8741	11072714 (2195, 2196)			UNCLASSIFIED	1034
8742	79211939 (2203, 2204)			UNCLASSIFIED	1054
8743	80462955 (2207, 2208)			UNCLASSIFIED	1012, 1038
8744	37022190 (2233, 2234)			UNCLASSIFIED	1025
8745	29983151 (2249, 2250)			UNCLASSIFIED	1030
8746	79867231 (2253, 2254)			UNCLASSIFIED	1017
8747	94233377 (2255, 2256)			UNCLASSIFIED	1010
8748	37036298 (2259, 2260)			UNCLASSIFIED	1012
8749	80450850 (2277, 2278)			UNCLASSIFIED	1010, 1023, 1034, 1038, 1044
8750	46778058 (2281, 2282)			UNCLASSIFIED	1021

8751	17900523 (2285, 2286)			UNCLASSIFIED	1044
8752	78980663 (2287, 2288)			UNCLASSIFIED	1038
8753	80055983 (2289, 2290)			UNCLASSIFIED	1010
8754	79264444 (2299, 2300)			UNCLASSIFIED	1019, 1024, 1054
8755	66259302 (2305, 2306)			UNCLASSIFIED	1010
8756	65328854 (2317, 2318)			UNCLASSIFIED	1022
8757	57310342 (2325, 2326)			UNCLASSIFIED	1024
8758	34766244 (2327, 2328)			UNCLASSIFIED	1050
8759	78517821 (2333, 2334)			UNCLASSIFIED	1003
8760	79329317 (2335, 2336)			UNCLASSIFIED	1053
8761	78706396 (2337, 2338)			UNCLASSIFIED	1008, 1026
8762	78463619 (2343, 2344)			UNCLASSIFIED	1039
8763	54538853 (2347, 2348)			UNCLASSIFIED	1040
8764	28119406 (2361, 2362)			UNCLASSIFIED	1016
8765	78467452 (2363, 2364)			UNCLASSIFIED	1026
8766	79250623 (2365, 2366)			UNCLASSIFIED	1008, 1022, 1023, 1024, 1026, 1054
8767	10344445 (2371, 2372)			UNCLASSIFIED	1038
8768	56931739 (2373, 2374)			UNCLASSIFIED	1025, 1054

8769	29647720 (2379, 2380)			UNCLASSIFIED	1022
8770	80580379 (2393, 2394)			UNCLASSIFIED	1038, 1058
8771	13893448 (2401, 2402)			UNCLASSIFIED	1034
8772	25327776 (2405, 2406)			UNCLASSIFIED	1008
8773	15001343 (2421, 2422)			UNCLASSIFIED	1027
8774	78367641 (2427, 2428)			UNCLASSIFIED	1003
8775	82361280 (2435, 2436)			UNCLASSIFIED	1001
8776	80027716 (2441, 2442)			UNCLASSIFIED	1001, 1004, 1024, 1025, 1044, 1054
8777	36997202 (2469, 2470)			UNCLASSIFIED	1001
8778	78769283 (2497, 2498)			UNCLASSIFIED	1008
8779	29191768 (2501, 2502)			UNCLASSIFIED	1003, 1008, 1017, 1023, 1039
8780	79261826 (2503, 2504)			UNCLASSIFIED	1023
8781	54994269 (2509, 2510)			UNCLASSIFIED	1029
8782	77868425 (2519, 2520)			UNCLASSIFIED	1041
8783	32746618 (2527, 2528)			UNCLASSIFIED	1000
8784	79586692 (2531, 2532)			UNCLASSIFIED	1001
8785	20294221 (2547, 2548)			UNCLASSIFIED	1022
8786	23283768 (2555, 2556)			UNCLASSIFIED	1003

8787	65704138 (2573, 2574)			UNCLASSIFIED	1038
8788	65657552 (2577, 2578)			UNCLASSIFIED	1010
8789	29916355 (2591, 2592)			UNCLASSIFIED	1030
8790	10330591 (2593, 2594)			UNCLASSIFIED	1025
8791	86468870 (2603, 2604)			UNCLASSIFIED	1029
8792	11770840 (2615, 2616)			UNCLASSIFIED	1025
8793	8491167 (2617, 2618)			UNCLASSIFIED	1034
8794	37799103 (2621, 2622)			UNCLASSIFIED	1012
8795	27963733 (2623, 2624)			UNCLASSIFIED	1022
8796	16373538 (2627, 2628)			UNCLASSIFIED	1030
8797	79858316 (2628, 2630)			UNCLASSIFIED	1003, 1019, 1038
8798	20634720 (2633, 2634)			UNCLASSIFIED	1022
8799	27367160 (2637, 2638)			UNCLASSIFIED	1022
8800	34623791 (2643, 2644)			UNCLASSIFIED	1029
8801	20371165 (2645, 2646)			UNCLASSIFIED	1022
8802	65702496 (2649, 2650)			UNCLASSIFIED	1038
8803	56066159 (2651, 2652)			UNCLASSIFIED	1021
8804	79910127 (2653, 2654)			UNCLASSIFIED	1038, 1044

8805	79556482 (2667, 2668)			UNCLASSIFIED	1039
8806	77709997 (2675, 2676)			UNCLASSIFIED	1054
8807	20467239 (2681, 2682)			UNCLASSIFIED	1010
8808	87454832 (2683, 2684)			UNCLASSIFIED	1026
8809	25328101 (2695, 2696)			UNCLASSIFIED	1044
8810	10223337 (2701, 2702)			UNCLASSIFIED	1006
8811	33099401 (2703, 2704)			UNCLASSIFIED	1026
8812	65692076 (2707, 2708)			UNCLASSIFIED	1009
8813	49551490 (2725, 2726)			UNCLASSIFIED	1040
8814	78736624 (2731, 2732)			UNCLASSIFIED	1044
8815	30811956 (2735, 2736)			UNCLASSIFIED	1026
8816	28774236 (2737, 2738)			UNCLASSIFIED	1009, 1023
8817	25250984 (2753, 2754)			UNCLASSIFIED	1008
8818	20282039 (2767, 2768)			UNCLASSIFIED	1022
8819	95012623 (2767, 2768)			UNCLASSIFIED	1003
8820	10168386 (2771, 2772)			UNCLASSIFIED	1003
8821	20438593 (2779, 2780)			UNCLASSIFIED	1010
8822	54530799 (2799, 2800)			UNCLASSIFIED	1041

8823	51625936 (2809, 2810)			UNCLASSIFIED	1040
8824	19511295 (2811, 2812)			UNCLASSIFIED	1034
8825	95293083 (2817, 2818)			UNCLASSIFIED	1022, 1030
8826	28473469 (2827, 2828)			UNCLASSIFIED	1030
8827	36507304 (2829, 2830)			UNCLASSIFIED	1026, 1034
8828	32126018 (2831, 2832)			UNCLASSIFIED	1025, 1044
8829	78756246 (2835, 2836)			UNCLASSIFIED	1039
8830	80221136 (2839, 2840)			UNCLASSIFIED	1004, 1008, 1017, 1022, 1024, 1025, 1034, 1038
8831	13516434 (2843, 2844)			UNCLASSIFIED	1024
8832	9875136 (2851, 2852)			UNCLASSIFIED	1008
8833	86473010 (2859, 2860)			UNCLASSIFIED	1029
8834	65963700 (2869, 2870)			UNCLASSIFIED	1004
8835	65985985 (2881, 2882)			UNCLASSIFIED	1054
8836	78742147 (2883, 2884)			UNCLASSIFIED	1017
8837	78965419 (2893, 2894)			UNCLASSIFIED	1030
8838	78946330 (2923, 2924)			UNCLASSIFIED	1017
8839	66441283 (2933, 2934)			UNCLASSIFIED	1022

8840	86473385 (2939, 2940)			UNCLASSIFIED	1029
8841	27981564 (2947, 2948)			UNCLASSIFIED	1006
8842	85800864 (2957, 2958)			UNCLASSIFIED	1025, 1029
8843	94848069 (2975, 2976)			UNCLASSIFIED	1044
8844	77969612 (2981, 2982)			UNCLASSIFIED	1040
8845	8740377 (2983, 2984)			UNCLASSIFIED	1022
8846	33202378 (2993, 2994)			UNCLASSIFIED	1026
8847	91236530 (2997, 2998)			UNCLASSIFIED	1010
8848	52563739 (3001, 3002)			UNCLASSIFIED	1044
8849	17905872 (3005, 3006)			UNCLASSIFIED	1006, 1044
8850	20612392 (3011, 3012)			UNCLASSIFIED	1010
8851	65468614 (3013, 3014)			UNCLASSIFIED	1054
8852	79631888 (3015, 3016)			UNCLASSIFIED	1024, 1027, 1044
8853	66396141 (3035, 3036)			UNCLASSIFIED	1008
8854	77695353 (3039, 3040)			UNCLASSIFIED	1044
8855	87461948 (3051, 3052)			UNCLASSIFIED	1015
8856	79841933 (3059, 3060)			UNCLASSIFIED	1044
8857	36990819 (3063, 3064)			UNCLASSIFIED	1016

8858	6510001 (3065, 3066)			UNCLASSIFIED	1022
8859	78468703 (3071, 3072)			UNCLASSIFIED	1026
8860	79163959 (3077, 3078)			UNCLASSIFIED	1004
8861	79329023 (3081, 3082)			UNCLASSIFIED	1053
8862	37448189 (3085, 3086)			UNCLASSIFIED	1034
8863	77516524 (3103, 3104)			UNCLASSIFIED	1049
8864	78406753 (3107, 3108)			UNCLASSIFIED	1024
8865	25316451 (3109, 3110)			UNCLASSIFIED	1009
8866	79115716 (3123, 3124)			UNCLASSIFIED	1016, 1024, 1044
8867	27842624 (3135, 3136)			UNCLASSIFIED	1034
8868	87447545 (3145, 3146)			UNCLASSIFIED	1025, 1029
8869	9846646 (3147, 3148)			UNCLASSIFIED	1023
8870	65698400 (3159, 3160)			UNCLASSIFIED	1012, 1038
8871	78940687 (3167, 3168)			UNCLASSIFIED	1026, 1039
8872	30629622 (3169, 3170)			UNCLASSIFIED	1026
8873	79568340 (3173, 3174)			UNCLASSIFIED	1003, 1017
8874	17290762 (3175, 3176)			UNCLASSIFIED	1054
8875	38438948 (3181, 3182)			UNCLASSIFIED	1034

8876	19520531 (3189, 3190)			UNCLASSIFIED	1025
8877	28775109 (3195, 3196)			UNCLASSIFIED	1009
8878	19637567 (3199, 3200)			UNCLASSIFIED	1054
8879	78317397 (3205, 3206)			UNCLASSIFIED	1026
8880	10325877 (3219, 3220)			UNCLASSIFIED	1025
8881	65663321 (3223, 3224)			UNCLASSIFIED	1053
8882	78793521 (3225, 3226)			UNCLASSIFIED	1009
8883	78963776 (3227, 3228)			UNCLASSIFIED	1003, 1008
8884	79896979 (3229, 3230)			UNCLASSIFIED	1017
8885	79255576 (3235, 3236)			UNCLASSIFIED	1008, 1023
8886	91009129 (3245, 3246)			UNCLASSIFIED	1010
8887	91228515 (3247, 3248)			UNCLASSIFIED	1010, 1040
8888	35068432 (3249, 3250)			UNCLASSIFIED	1030
8889	78941357 (3291, 3292)			UNCLASSIFIED	1039
8890	30257689 (3303, 3304)			UNCLASSIFIED	1026
8891	35142221 (3305, 3306)			UNCLASSIFIED	1050
8892	55186431 (3333, 3334)			UNCLASSIFIED	1019
8893	78254399 (3343, 3344)			UNCLASSIFIED	1026

8894	78491386 (3349, 3350)			UNCLASSIFIED	1054
8895	77712326 (3351, 3352)			UNCLASSIFIED	1054
8896	79115180 (3359, 3360)			UNCLASSIFIED	1003
8897	56065583 (3361, 3362)			UNCLASSIFIED	1042
8898	14403161 (3369, 3370)			UNCLASSIFIED	1053
8899	79491945 (3375, 3376)			UNCLASSIFIED	1038, 1053
8900	55449691 (3377, 3378)			UNCLASSIFIED	1009
8901	80191220 (3379, 3380)			UNCLASSIFIED	1034, 1058
8902	30631902 (3387, 3388)			UNCLASSIFIED	1022
8903	46574257 (3389, 3390)			UNCLASSIFIED	1054
8904	30537342 (3391, 3392)			UNCLASSIFIED	1026
8905	13076100 (3395, 3396)			UNCLASSIFIED	1027
8906	65704328 (3397, 3398)			UNCLASSIFIED	1023
8907	10885643 (3409, 3410)			UNCLASSIFIED	1031
8908	80086169 (3411, 3412)			UNCLASSIFIED	1024
8909	78491890 (3427, 3428)			UNCLASSIFIED	1054
8910	86694625 (3451, 3452)			UNCLASSIFIED	1018, 1029
8911	20474349 (3457, 3458)			UNCLASSIFIED	1022

8912	69960717 (3459, 3460)			UNCLASSIFIED	1042
8913	65671346 (3475, 3476)			UNCLASSIFIED	1016
8914	54698643 (3477, 3478)			UNCLASSIFIED	1041
8915	15024349 (3483, 3484)			UNCLASSIFIED	1054
8916	27213508 (3485, 3486)			UNCLASSIFIED	1039
8917	66397011 (3493, 3494)			UNCLASSIFIED	1009
8918	70429617 (3503, 3504)			UNCLASSIFIED	1019, 1041
8919	11775754 (3507, 3508)			UNCLASSIFIED	1031
8920	80088083 (3513, 3514)			UNCLASSIFIED	1004, 1039
8921	70960193 (3515, 3516)			UNCLASSIFIED	1029
8922	20205916 (3517, 3518)			UNCLASSIFIED	1010
8923	78051854 (3519, 3520)			UNCLASSIFIED	1026
8924	30198003 (3521, 3522)			UNCLASSIFIED	1026
8925	78194517 (3527, 3528)			UNCLASSIFIED	1034, 1049
8926	29020604 (3539, 3540)			UNCLASSIFIED	1008
8927	79778305 (3547, 3548)			UNCLASSIFIED	1003, 1017
8928	66429997 (3557, 3558)			UNCLASSIFIED	1024
8929	8022712 (3561, 3562)			UNCLASSIFIED	1022

8930	38901541 (3571, 3572)			UNCLASSIFIED	1024
8931	56072722 (3573, 3574)			UNCLASSIFIED	1021
8932	10144093 (3585, 3586)			UNCLASSIFIED	1001
8933	66436786 (3605, 3606)			UNCLASSIFIED	1024
8934	20432737 (3607, 3608)			UNCLASSIFIED	1024
8935	87451852 (3633, 3634)			UNCLASSIFIED	1025
8936	79571951 (3635, 3636)			UNCLASSIFIED	1017
8937	28361885 (3647, 3648)			UNCLASSIFIED	1024
8938	65882846 (3655, 3656)			UNCLASSIFIED	1054
8939	24128902 (3661, 3662)			UNCLASSIFIED	1009
8940	77530773 (3681, 3682)			UNCLASSIFIED	1038
8941	94663581 (3683, 3684)			UNCLASSIFIED	1016, 1023, 1024, 1054
8942	17938806 (3691, 3692)			UNCLASSIFIED	1038
8943	91256285 (3701, 3702)			UNCLASSIFIED	1010
8944	71770206 (3705, 3706)			UNCLASSIFIED	1003, 1017
8945	65648227 (3709, 3710)			UNCLASSIFIED	1054
8946	79629729 (3717, 3718)			UNCLASSIFIED	1039
8947	29024681 (3731, 3732)			UNCLASSIFIED	1026

8948	23332512 (3735, 3736)			UNCLASSIFIED	1053
8949	78300815 (3737, 3738)			UNCLASSIFIED	1026
8950	10093080 (3739, 3740)			UNCLASSIFIED	1038
8951	15032155 (3745, 3746)			UNCLASSIFIED	1054
8952	32714734 (3749, 3750)			UNCLASSIFIED	1029
8953	52234444 (3751, 3752)			UNCLASSIFIED	1036
8954	87100424 (3753, 3754)			UNCLASSIFIED	1022, 1024
8955	86474604 (3763, 3764)			UNCLASSIFIED	1029
8956	33204297 (3767, 3768)			UNCLASSIFIED	1026
8957	17956632 (3769, 3770)			UNCLASSIFIED	1025
8958	78365815 (3773, 3774)			UNCLASSIFIED	1026
8959	94147448 (3779, 3780)			UNCLASSIFIED	1003, 1010, 1044
8960	79190311 (3787, 3788)			UNCLASSIFIED	1024
8961	10119455 (3797, 3798)			UNCLASSIFIED	1023
8962	29917341 (3813, 3814)			UNCLASSIFIED	1022
8963	80054123 (3819, 3820)			UNCLASSIFIED	1001, 1022
8964	87897508 (3821, 3822)			UNCLASSIFIED	1015
8965	20484787 (3823, 3824)			UNCLASSIFIED	1010

8966	11606794 (3827, 3828)			UNCLASSIFIED	1024
8967	47652402 (3841, 3842)			UNCLASSIFIED	1029
8968	11394886 (3855, 3856)			UNCLASSIFIED	1010
8969	20297507 (3857, 3858)			UNCLASSIFIED	1034
8970	11306148 (3863, 3864)			UNCLASSIFIED	1022
8971	46868814 (3865, 3866)			UNCLASSIFIED	1034
8972	28802403 (3875, 3876)			UNCLASSIFIED	1022
8973	46868258 (3879, 3880)			UNCLASSIFIED	1037
8974	20695452 (3897, 3898)			UNCLASSIFIED	1022
8975	78996159 (3903, 3904)			UNCLASSIFIED	1008, 1017
8976	16504576 (3905, 3906)			UNCLASSIFIED	1008
8977	28097888 (3913, 3914)			UNCLASSIFIED	1054
8978	30199928 (3927, 3928)			UNCLASSIFIED	1026
8979	8230680 (3939, 3940)			UNCLASSIFIED	1004
8980	27976902 (3957, 3958)			UNCLASSIFIED	1022
8981	65641514 (3979, 3980)			UNCLASSIFIED	1041
8982	82321406 (3995, 3996)			UNCLASSIFIED	1001, 1006, 1022, 1029, 1054
8983	86691481 (4005, 4006)			UNCLASSIFIED	1018

8984	17735043 (4013, 4014)			UNCLASSIFIED	1003
8985	28954757 (4025, 4026)			UNCLASSIFIED	1022
8986	78387002 (4029, 4030)			UNCLASSIFIED	1026
8987	56071133 (4033, 4034)			UNCLASSIFIED	1010
8988	66403949 (4035, 4036)			UNCLASSIFIED	1022
8989	30454354 (4037, 4038)			UNCLASSIFIED	1026
8990	78719491 (4041, 4042)			UNCLASSIFIED	1008, 1009, 1022, 1023, 1026
8991	50400915 (4043, 4044)			UNCLASSIFIED	1036
8992	20290968 (4047, 4048)			UNCLASSIFIED	1022
8993	46868608 (4053, 4054)			UNCLASSIFIED	1037
8994	32435065 (4069, 4070)			UNCLASSIFIED	1029
8995	29456185 (4077, 4078)			UNCLASSIFIED	1022
8996	81903046 (4079, 4080)			UNCLASSIFIED	1054
8997	77918506 (4089, 4090)			UNCLASSIFIED	1036
8998	66467906 (4101, 4102)			UNCLASSIFIED	1001, 1024
8999	18600143 (4111, 4112)			UNCLASSIFIED	1023
9000	17882559 (4113, 4114)			UNCLASSIFIED	1024
9001	42723985 (4125, 4126)			UNCLASSIFIED	1050

9002	87466764 (4131, 4132)			UNCLASSIFIED	1015
9003	79856553 (4137, 4138)			UNCLASSIFIED	1017
9004	80034616 (4155, 4156)			UNCLASSIFIED	1053
9005	11817460 (4165, 4166)			UNCLASSIFIED	1006
9006	77877721 (4169, 4170)			UNCLASSIFIED	1019
9007	35070476 (4173, 4174)			UNCLASSIFIED	1030
9008	39565458 (4197, 4198)			UNCLASSIFIED	1024
9009	27978124 (4199, 4200)			UNCLASSIFIED	1022
9010	65658588 (4217, 4218)			UNCLASSIFIED	1054
9011	57567558 (4219, 4220)			UNCLASSIFIED	1023
9012	10063472 (4221, 4222)			UNCLASSIFIED	1017
9013	65985994 (4241, 4242)			UNCLASSIFIED	1054
9014	21659614 (4249, 4250)			UNCLASSIFIED	1034
9015	79423490 (4261, 4262)			UNCLASSIFIED	1017
9016	79202903 (4263, 4264)			UNCLASSIFIED	1001
9017	80254010 (4279, 4280)			UNCLASSIFIED	1054
9018	21150781 (4281, 4282)			UNCLASSIFIED	1024
9019	66258577 (4285, 4286)			UNCLASSIFIED	1010, 1023

9020	27968300 (4295, 4296)			UNCLASSIFIED	1006
9021	55147667 (4299, 4300)			UNCLASSIFIED	1034
9022	28487635 (4303, 4304)			UNCLASSIFIED	1039
9023	79959997 (4309, 4310)			UNCLASSIFIED	1008, 1036
9024	77781590 (4311, 4312)			UNCLASSIFIED	1019
9025	37811518 (4321, 4322)			UNCLASSIFIED	1058
9026	29910570 (4341, 4342)			UNCLASSIFIED	1030
9027	65452245 (4343, 4344)			UNCLASSIFIED	1016, 1038
9028	79642599 (4347, 4348)			UNCLASSIFIED	1022, 1038
9029	36521449 (4353, 4354)			UNCLASSIFIED	1029
9030	71085452 (4365, 4366)			UNCLASSIFIED	1016, 1037
9031	65680814 (4387, 4388)			UNCLASSIFIED	1041
9032	85516344 (4389, 4390)			UNCLASSIFIED	1049
9033	17298534 (4397, 4398)			UNCLASSIFIED	1023
9034	65874208 (4413, 4414)			UNCLASSIFIED	1016
9035	49123340 (4433, 4434)			UNCLASSIFIED	1040
9036	25320510 (4439, 4440)			UNCLASSIFIED	1003
9037	78967306 (4441, 4442)			UNCLASSIFIED	1050, 1054

9038	78926378 (4445, 4446)			UNCLASSIFIED	1039
9039	79599993 (4447, 4448)			UNCLASSIFIED	1034
9040	87894813 (4457, 4458)			UNCLASSIFIED	1018
9041	33181789 (4461, 4462)			UNCLASSIFIED	1026
9042	54509725 (4479, 4480)			UNCLASSIFIED	1042
9043	80047982 (4485, 4486)			UNCLASSIFIED	1024
9044	65893545 (4487, 4488)			UNCLASSIFIED	1038
9045	16836327 (4493, 4494)			UNCLASSIFIED	1003
9046	20112811 (4495, 4496)			UNCLASSIFIED	1034
9047	56134052 (4501, 4502)			UNCLASSIFIED	1026
9048	78500141 (4519, 4520)			UNCLASSIFIED	1022
9049	78721695 (4531, 4532)			UNCLASSIFIED	1050, 1054
9050	35017420 (4533, 4534)			UNCLASSIFIED	1054
9051	27366845 (4543, 4544)			UNCLASSIFIED	1026
9052	98198505 (4545, 4546)			UNCLASSIFIED	1012, 1034, 1053
9053	27963217 (4555, 4556)			UNCLASSIFIED	1022
9054	28470928 (4557, 4558)			UNCLASSIFIED	1054
9055	29014585 (4573, 4574)			UNCLASSIFIED	1044

9056	37009991 (4577, 4578)			UNCLASSIFIED	1029
9057	11100641 (4587, 4588)			UNCLASSIFIED	1022
9058	79560541 (4591, 4592)			UNCLASSIFIED	1038
9059	65671769 (4593, 4594)			UNCLASSIFIED	1016
9060	6619451 (4601, 4602)			UNCLASSIFIED	1022
9061	77911777 (4611, 4612)			UNCLASSIFIED	1036
9062	65677759 (4619, 4620)			UNCLASSIFIED	1016
9063	87114165 (4621, 4622)			UNCLASSIFIED	1025
9064	10131031 (4623, 4624)			UNCLASSIFIED	1016, 1029, 1039
9065	65660942 (4625, 4626)			UNCLASSIFIED	1024
9066	65857768 (4627, 4628)			UNCLASSIFIED	1016, 1038
9067	39372286 (4629, 4630)			UNCLASSIFIED	1022, 1058
9068	78786236 (4631, 4632)			UNCLASSIFIED	1022
9069	27832114 (4635, 4636)			UNCLASSIFIED	1004, 1022, 1024, 1039
9070	10248589 (4637, 4638)			UNCLASSIFIED	1039
9071	78474297 (4651, 4652)			UNCLASSIFIED	1003, 1044
9072	46856553 (4657, 4658)			UNCLASSIFIED	1029
9073	78898942 (4661, 4662)			UNCLASSIFIED	1017

9074	57052092 (4663, 4664)			UNCLASSIFIED	1049
9075	77692114 (4665, 4666)			UNCLASSIFIED	1024
9076	46770016 (4677, 4678)			UNCLASSIFIED	1041
9077	65446930 (4687, 4688)			UNCLASSIFIED	1038
9078	13506991 (4701, 4702)			UNCLASSIFIED	1027
9079	9857038 (4703, 4704)			UNCLASSIFIED	1008
9080	94112356 (4709, 4710)			UNCLASSIFIED	1016, 1034, 1037, 1054
9081	78459973 (4715, 4716)			UNCLASSIFIED	1026
9082	29352843 (4721, 4722)			UNCLASSIFIED	1022
9083	10029658 (4727, 4728)			UNCLASSIFIED	1039
9084	10194259 (4729, 4730)			UNCLASSIFIED	1003
9085	94846426 (4753, 4754)			UNCLASSIFIED	1044
9086	78259273 (4757, 4758)			UNCLASSIFIED	1026
9087	79239699 (4765, 4766)			UNCLASSIFIED	1003
9088	80502530 (4769, 4770)			UNCLASSIFIED	1012
9089	79320132 (4773, 4774)			UNCLASSIFIED	1022, 1037
9090	52563626 (4785, 4786)			UNCLASSIFIED	1044
9091	79844475 (4797, 4798)			UNCLASSIFIED	1017

9092	80503996 (4835, 4836)			UNCLASSIFIED	1004, 1012, 1022, 1024, 1031, 1034
9093	10155588 (4839, 4840)			UNCLASSIFIED	1039
9094	36830219 (4857, 4858)			UNCLASSIFIED	1054
9095	82335125 (4865, 4866)			UNCLASSIFIED	1001, 1024, 1038, 1058
9096	87112263 (4871, 4872)			UNCLASSIFIED	1007
9097	5829683 (4881, 4882)			UNCLASSIFIED	1058
9098	56516521 (4889, 4890)			UNCLASSIFIED	1019
9099	66392546 (4895, 4896)			UNCLASSIFIED	1003
9100	16312877 (4907, 4908)			UNCLASSIFIED	1024
9101	10370679 (4909, 4910)			UNCLASSIFIED	1044
9102	77572804 (4921, 4922)			UNCLASSIFIED	1023
9103	78488338 (4947, 4948)			UNCLASSIFIED	1026
9104	79261045 (4949, 4950)			UNCLASSIFIED	1024
9105	11809865 (4951, 4952)			UNCLASSIFIED	1022
9106	46849908 (4959, 4960)			UNCLASSIFIED	1029
9107	36511971 (4961, 4962)			UNCLASSIFIED	1026
9108	80408013 (4973, 4974)			UNCLASSIFIED	1003, 1029
9109	33198917 (4981, 4982)			UNCLASSIFIED	1029

9110	13521592 (5005, 5006)			UNCLASSIFIED	1024
9111	25258958 (5007, 5008)			UNCLASSIFIED	1003
9112	11817088 (5013, 5014)			UNCLASSIFIED	1006
9113	82418377 (5017, 5018)			UNCLASSIFIED	1001, 1006, 1010, 1012, 1022, 1044, 1068
9114	80232437 (5027, 5028)			UNCLASSIFIED	1001, 1003, 1004, 1008, 1010, 1017, 1024, 1039, 1046, 1054
9115	39517743 (5033, 5034)			UNCLASSIFIED	1006, 1034
9116	11687042 (5043, 5044)			UNCLASSIFIED	1031
9117	11695136 (5045, 5046)			UNCLASSIFIED	1034
9118	16380963 (5047, 5048)			UNCLASSIFIED	1030
9119	25144960 (5055, 5056)			UNCLASSIFIED	1017
9120	46692673 (5057, 5058)			UNCLASSIFIED	1034
9121	78890669 (5067, 5068)			UNCLASSIFIED	1044
9122	56155227 (5071, 5072)			UNCLASSIFIED	1021
9123	37420764 (5083, 5084)			UNCLASSIFIED	1016
9124	21428141 (5089, 5090)			UNCLASSIFIED	1022
9125	87469140 (5123, 5124)			UNCLASSIFIED	1018

9126	11200994 (5129, 5130)			UNCLASSIFIED	1022
9127	78519122 (5133, 5134)			UNCLASSIFIED	1003
9128	27982165 (5141, 5142)			UNCLASSIFIED	1022
9129	29453540 (5143, 5144)			UNCLASSIFIED	1022
9130	34900016 (5169, 5170)			UNCLASSIFIED	1054
9131	33187094 (5175, 5176)			UNCLASSIFIED	1026
9132	86674854 (5183, 5184)			UNCLASSIFIED	1007
9133	32717912 (5197, 5198)			UNCLASSIFIED	1029
9134	28396226 (5201, 5202)			UNCLASSIFIED	1038
9135	94998851 (5203, 5204)			UNCLASSIFIED	1004
9136	78502028 (5233, 5234)			UNCLASSIFIED	1026
9137	79590398 (5243, 5244)			UNCLASSIFIED	1024, 1026
9138	39733149 (5247, 5248)			UNCLASSIFIED	1026
9139	32429948 (5261, 5262)			UNCLASSIFIED	1050
9140	80389847 (5267, 5268)			UNCLASSIFIED	1025, 1029
9141	28800992 (5269, 5270)			UNCLASSIFIED	1022
9142	48048571 (5277, 5278)			UNCLASSIFIED	1026, 1038, 1041
9143	79190835 (5279, 5280)			UNCLASSIFIED	1004

9144	65879823 (5283, 5284)			UNCLASSIFIED	1000, 1006
9145	86380260 (5297, 5298)			UNCLASSIFIED	1053
9146	80048399 (5311, 5312)			UNCLASSIFIED	1006, 1008, 1010
9147	27962552 (5313, 5314)			UNCLASSIFIED	1022
9148	65501060 (5315, 5316)			UNCLASSIFIED	1016
9149	49529902 (5317, 5318)			UNCLASSIFIED	1036
9150	80306968 (5323, 5324)			UNCLASSIFIED	1008, 1022, 1029
9151	10801108 (5333, 5334)			UNCLASSIFIED	1003
9152	30485755 (5341, 5342)			UNCLASSIFIED	1030, 1044
9153	80248348 (5345, 5346)			UNCLASSIFIED	1003, 1004, 1006, 1016, 1017, 1031, 1034, 1039, 1054
9154	79844225 (5351, 5352)			UNCLASSIFIED	1008
9155	85805347 (5359, 5360)			UNCLASSIFIED	1054
9156	21641890 (5363, 5364)			UNCLASSIFIED	1022
9157	79199356 (5365, 5366)			UNCLASSIFIED	1054
9158	19635276 (5369, 5370)			UNCLASSIFIED	1024
9159	34394173 (5371, 5372)			UNCLASSIFIED	1026
9160	54527923 (5379, 5380)			UNCLASSIFIED	1041

9161	77881592 (5387, 5388)			UNCLASSIFIED	1019
9162	95288391 (5389, 5390)			UNCLASSIFIED	1004, 1017, 1024, 1038
9163	65880574 (5397, 5398)			UNCLASSIFIED	1054
9164	65506470 (5401, 5402)			UNCLASSIFIED	1016
9165	80054377 (5403, 5404)			UNCLASSIFIED	1022
9166	33188378 (5417, 5418)			UNCLASSIFIED	1026
9167	79814400 (5419, 5420)			UNCLASSIFIED	1017
9168	19881910 (5431, 5432)			UNCLASSIFIED	1034
9169	78678243 (5433, 5434)			UNCLASSIFIED	1028
9170	71844116 (5443, 5444)			UNCLASSIFIED	1022
9171	66001061 (5445, 5446)			UNCLASSIFIED	1006
9172	28953323 (5463, 5464)			UNCLASSIFIED	1003
9173	78163702 (5469, 5470)			UNCLASSIFIED	1024, 1029
9174	20724574 (5471, 5472)			UNCLASSIFIED	1004
9175	10237679 (5489, 5490)			UNCLASSIFIED	1025
9176	79968361 (5493, 5494)			UNCLASSIFIED	1025, 1039
9177	57050847 (5497, 5498)			UNCLASSIFIED	1023
9178	27825616 (5507, 5508)			UNCLASSIFIED	1053

9179	9358682 (5513, 5514)			UNCLASSIFIED	1017
9180	52385412 (5515, 5516)			UNCLASSIFIED	1036
9181	39725383 (5521, 5522)			UNCLASSIFIED	1054
9182	86476530 (5525, 5526)			UNCLASSIFIED	1029
9183	20370378 (5535, 5536)			UNCLASSIFIED	1010
9184	13016243 (5543, 5544)			UNCLASSIFIED	1004
9185	77554298 (5545, 5546)			UNCLASSIFIED	1054
9186	54539627 (5557, 5558)			UNCLASSIFIED	1041
9187	79333747 (5559, 5560)			UNCLASSIFIED	1008
9188	10853737 (5563, 5564)			UNCLASSIFIED	1017
9189	23332997 (5567, 5568)			UNCLASSIFIED	1044
9190	80359544 (5591, 5592)			UNCLASSIFIED	1008, 1017, 1023, 1024, 1029
9191	28483422 (5595, 5596)			UNCLASSIFIED	1017
9192	65693538 (5601, 5602)			UNCLASSIFIED	1009
9193	48042587 (5611, 5612)			UNCLASSIFIED	1042
9194	46899294 (5619, 5620)			UNCLASSIFIED	1041
9195	66001063 (5627, 5628)			UNCLASSIFIED	1006, 1016, 1058
9196	85807649 (5629, 5630)			UNCLASSIFIED	1019, 1029, 1054

9197	66762313 (5641, 5642)			UNCLASSIFIED	1024
9198	86380029 (5647, 5648)			UNCLASSIFIED	1053
9199	86380266 (5649, 5650)			UNCLASSIFIED	1053
9200	19464159 (5651, 5652)			UNCLASSIFIED	1039
9201	79462878 (5653, 5654)			UNCLASSIFIED	1031, 1038
9202	10300869 (5655, 5656)			UNCLASSIFIED	1025, 1038
9203	80074244 (5661, 5662)			UNCLASSIFIED	1022
9204	82059644 (5675, 5676)			UNCLASSIFIED	1006, 1016, 1034
9205	85801837 (5681, 5682)			UNCLASSIFIED	1038
9206	25260565 (5685, 5686)			UNCLASSIFIED	1008
9207	66435404 (5721, 5722)			UNCLASSIFIED	1024
9208	79557566 (5723, 5724)			UNCLASSIFIED	1025
9209	86475994 (5731, 5732)			UNCLASSIFIED	1029
9210	27846192 (5733, 5734)			UNCLASSIFIED	1004, 1022, 1034
9211	51578522 (5739, 5740)			UNCLASSIFIED	1036
9212	79827801 (5741, 5742)			UNCLASSIFIED	1003
9213	80090131 (5749, 5750)			UNCLASSIFIED	1008, 1024
9214	12882651 (5751, 5752)			UNCLASSIFIED	1004

9215	20705925 (5769, 5770)			UNCLASSIFIED	1022
9216	47656179 (5779, 5780)			UNCLASSIFIED	1029
9217	78246389 (5783, 5784)			UNCLASSIFIED	1008
9218	38348808 (5785, 5786)			UNCLASSIFIED	1026
9219	79563436 (5801, 5802)			UNCLASSIFIED	1001
9220	78906817 (5841, 5842)			UNCLASSIFIED	1023
9221	30385688 (5843, 5844)			UNCLASSIFIED	1016
9222	88094271 (5847, 5848)			UNCLASSIFIED	1034
9223	78383565 (5851, 5852)			UNCLASSIFIED	1003
9224	98198669 (5871, 5872)			UNCLASSIFIED	1004, 1006, 1012, 1016, 1019, 1023, 1026
9225	35606160 (5879, 5880)			UNCLASSIFIED	1053
9226	5806994 (5885, 5886)			UNCLASSIFIED	1058
9227	8485932 (5901, 5902)			UNCLASSIFIED	1010
9228	11102556 (5907, 5908)			UNCLASSIFIED	1053
9229	78939342 (5909, 5910)			UNCLASSIFIED	1023, 1026
9230	94148498 (5911, 5912)			UNCLASSIFIED	1003
9231	13088430 (5945, 5946)			UNCLASSIFIED	1027
9232	78725270 (5947, 5948)			UNCLASSIFIED	1008

9233	46805633 (5963, 5954)			UNCLASSIFIED	1034
9234	56660043 (5955, 5956)			UNCLASSIFIED	1042
9235	38863326 (5961, 5962)			UNCLASSIFIED	1022
9236	30650854 (5965, 5966)			UNCLASSIFIED	1008
9237	86684669 (5969, 5970)			UNCLASSIFIED	1029, 1054
9238	79176430 (5973, 5974)			UNCLASSIFIED	1004, 1034, 1054, 1055
9239	82238402 (5981, 5982)			UNCLASSIFIED	1016
9240	28329651 (5989, 5990)			UNCLASSIFIED	1024
9241	36990837 (5993, 5994)			UNCLASSIFIED	1016
9242	51627406 (6001, 6002)			UNCLASSIFIED	1019
9243	65655733 (6003, 6004)			UNCLASSIFIED	1054
9244	32746630 (6041, 6042)			UNCLASSIFIED	1000
9245	10856067 (6045, 6046)			UNCLASSIFIED	1001
9246	79959681 (6049, 6050)			UNCLASSIFIED	1034
9247	29688125 (6053, 6054)			UNCLASSIFIED	1024
9248	80583900 (6057, 6058)			UNCLASSIFIED	1008, 1058
9249	79205039 (6059, 6060)			UNCLASSIFIED	1024
9250	65877878 (6065, 6066)			UNCLASSIFIED	1038

9251	11686230 (6067, 6068)			UNCLASSIFIED	1034
9252	65502750 (6075, 6076)			UNCLASSIFIED	1016
9253	78069107 (6081, 6082)			UNCLASSIFIED	1036, 1037
9254	35108317 (6085, 6086)			UNCLASSIFIED	1030
9255	21415106 (6095, 6096)			UNCLASSIFIED	1022
9256	65479512 (6099, 6100)			UNCLASSIFIED	1054
9257	11259145 (6117, 6118)			UNCLASSIFIED	1022
9258	91213986 (6131, 6132)			UNCLASSIFIED	1024
9259	8491185 (6143, 6144)			UNCLASSIFIED	1024
9260	65864995 (6167, 6168)			UNCLASSIFIED	1016
9261	38346549 (6183, 6184)			UNCLASSIFIED	1008
9262	78295056 (6187, 6188)			UNCLASSIFIED	1038
9263	88094914 (6199, 6200)			UNCLASSIFIED	1024
9264	19451962 (6203, 6204)			UNCLASSIFIED	1017
9265	79826770 (6207, 6208)			UNCLASSIFIED	1009, 1044
9266	29684848 (6209, 6210)			UNCLASSIFIED	1054
9267	30536784 (6217, 6218)			UNCLASSIFIED	1026
9268	30536958 (6223, 6224)			UNCLASSIFIED	1026

9269	57303796 (6231, 6232)			UNCLASSIFIED	1004, 1008, 1008, 1009, 1010, 1024, 1034
9270	9893524 (6235, 6236)			UNCLASSIFIED	1044
9271	11696061 (6241, 6242)			UNCLASSIFIED	1034
9272	79612675 (6247, 6248)			UNCLASSIFIED	1001, 1025, 1049, 1058
9273	35900868 (6253, 6254)			UNCLASSIFIED	1034
9274	80079782 (6261, 6262)			UNCLASSIFIED	1003, 1034
9275	16280326 (6263, 6264)			UNCLASSIFIED	1054
9276	20699394 (6273, 6274)			UNCLASSIFIED	1022
9277	33202293 (6291, 6292)			UNCLASSIFIED	1026
9278	55305796 (6295, 6296)			UNCLASSIFIED	1042
9279	78385367 (6305, 6306)			UNCLASSIFIED	1026
9280	20297638 (6307, 6308)			UNCLASSIFIED	1034
9281	30251698 (6321, 6322)			UNCLASSIFIED	1044
9282	36512814 (6329, 6330)			UNCLASSIFIED	1026
9283	39730685 (6339, 6340)			UNCLASSIFIED	1026
9284	78511145 (6343, 6344)			UNCLASSIFIED	1054
9285	95091988 (6349, 6350)			UNCLASSIFIED	1012
9286	55182043 (6361, 6362)			UNCLASSIFIED	1019

9287	94848435 (6375, 6376)			UNCLASSIFIED	1029, 1044
9288	65505003 (6395, 6396)			UNCLASSIFIED	1016
9289	25316110 (6397, 6398)			UNCLASSIFIED	1008
9290	78520201 (6399, 6400)			UNCLASSIFIED	1008
9291	10053155 (6417, 6418)			UNCLASSIFIED	1038
9292	78186519 (6421, 6422)			UNCLASSIFIED	1016, 1033, 1054, 1055
9293	65674955 (6437, 6438)			UNCLASSIFIED	1046
9294	80050214 (6439, 6440)			UNCLASSIFIED	1004, 1008
9295	79964343 (6441, 6442)			UNCLASSIFIED	1038
9296	30658121 (6449, 6450)			UNCLASSIFIED	1026
9297	86380895 (6453, 6454)			UNCLASSIFIED	1053
9298	78795396 (6457, 6458)			UNCLASSIFIED	1026
9299	86925663 (6459, 6460)			UNCLASSIFIED	1022
9300	17254457 (6461, 6462)			UNCLASSIFIED	1044
9301	78797776 (6463, 6464)			UNCLASSIFIED	1009
9302	20483055 (6469, 6470)			UNCLASSIFIED	1022
9303	33210736 (6471, 6472)			UNCLASSIFIED	1012
9304	71750829 (6473, 6474)			UNCLASSIFIED	1003, 1022

9305	10887692 (6475, 6476)			UNCLASSIFIED	1024
9306	78956106 (6488, 6490)			UNCLASSIFIED	1008
9307	35892950 (6497, 6498)			UNCLASSIFIED	1053
9308	57292620 (6507, 6508)			UNCLASSIFIED	1024, 1034
9309	66430167 (6521, 6522)			UNCLASSIFIED	1024
9310	65700660 (6523, 6524)			UNCLASSIFIED	1038
9311	87452757 (6537, 6538)			UNCLASSIFIED	1025
9312	78385458 (6555, 6556)			UNCLASSIFIED	1026, 1053
9313	27829496 (6557, 6558)			UNCLASSIFIED	1010
9314	65670538 (6561, 6562)			UNCLASSIFIED	1038
9315	66171652 (6563, 6564)			UNCLASSIFIED	1038
9316	65881948 (6567, 6568)			UNCLASSIFIED	1054
9317	29442699 (6577, 6578)			UNCLASSIFIED	1016
9318	20472612 (6583, 6584)			UNCLASSIFIED	1022
9319	85526490 (6585, 6586)			UNCLASSIFIED	1038
9320	77875487 (6591, 6592)			UNCLASSIFIED	1019
9321	77818545 (6595, 6596)			UNCLASSIFIED	1024
9322	27928448 (6599, 6600)			UNCLASSIFIED	1008

9323	35101425 (6605, 6606)			UNCLASSIFIED	1024
9324	27980057 (6609, 6610)			UNCLASSIFIED	1006
9325	66176154 (6613, 6614)			UNCLASSIFIED	1016
9326	10859941 (6621, 6622)			UNCLASSIFIED	1016
9327	17884479 (6637, 6638)			UNCLASSIFIED	1039
9328	79596345 (6655, 6656)			UNCLASSIFIED	1025
9329	8522416 (6661, 6662)			UNCLASSIFIED	1034
9330	78243781 (6669, 6670)			UNCLASSIFIED	1026
9331	78300832 (6683, 6684)			UNCLASSIFIED	1026
9332	80206505 (6685, 6686)			UNCLASSIFIED	1053
9333	79910335 (6695, 6696)			UNCLASSIFIED	1017
9334	71759512 (6711, 6712)			UNCLASSIFIED	1008
9335	38207960 (6715, 6716)			UNCLASSIFIED	1026
9336	65505366 (6723, 6724)			UNCLASSIFIED	1016
9337	86668842 (6745, 6746)			UNCLASSIFIED	1029
9338	32598621 (6755, 6756)			UNCLASSIFIED	1034
9339	80034634 (6767, 6768)			UNCLASSIFIED	1053
9340	46850030 (6787, 6788)			UNCLASSIFIED	1029

9341	80082947 (6811, 6812)			UNCLASSIFIED	1025, 1034, 1038, 1050, 1054
9342	17709868 (6813, 6814)			UNCLASSIFIED	1044
9343	81355940 (6819, 6820)			UNCLASSIFIED	1001, 1004, 1034, 1038, 1054
9344	80092981 (6823, 6824)			UNCLASSIFIED	1006
9345	78684936 (6825, 6826)			UNCLASSIFIED	1003, 1025
9346	10844920 (6827, 6828)			UNCLASSIFIED	1017
9347	35997373 (6829, 6830)			UNCLASSIFIED	1029
9348	55902351 (6835, 6836)			UNCLASSIFIED	1026
9349	37013441 (6837, 6838)			UNCLASSIFIED	1029
9350	35022103 (6839, 6840)			UNCLASSIFIED	1016
9351	11287984 (6847, 6848)			UNCLASSIFIED	1006
9352	66373013 (6855, 6856)			UNCLASSIFIED	1038
9353	65709220 (6863, 6864)			UNCLASSIFIED	1022, 1023, 1024
9354	16416386 (6871, 6872)			UNCLASSIFIED	1054
9355	80236246 (6887, 6888)			UNCLASSIFIED	1022, 1023
9356	10285603 (6905, 6906)			UNCLASSIFIED	1025
9357	80504802 (6907, 6908)			UNCLASSIFIED	1012
9358	46686838 (6923, 6924)			UNCLASSIFIED	1034

9359	11090238 (6925, 6926)			UNCLASSIFIED	1004
9360	30392968 (6929, 6930)			UNCLASSIFIED	1026
9361	25298464 (6931, 6932)			UNCLASSIFIED	1003
9362	36997558 (6933, 6934)			UNCLASSIFIED	1001
9363	79629905 (6937, 6938)			UNCLASSIFIED	1039
9364	65709045 (6941, 6942)			UNCLASSIFIED	1023
9365	5799757 (6945, 6946)			UNCLASSIFIED	1058
9366	30535052 (6947, 6948)			UNCLASSIFIED	1026
9367	39914966 (6955, 6956)			UNCLASSIFIED	1029
9368	86684314 (6985, 6986)			UNCLASSIFIED	1029
9369	19845111 (6987, 6988)			UNCLASSIFIED	1006
9370	35069096 (7001, 7002)			UNCLASSIFIED	1030
9371	79585168 (7009, 7010)			UNCLASSIFIED	1024
9372	71088760 (7011, 7012)			UNCLASSIFIED	1053
9373	80046750 (7019, 7020)			UNCLASSIFIED	1024
9374	17870227 (7021, 7022)			UNCLASSIFIED	1024
9375	86681838 (7023, 7024)			UNCLASSIFIED	1019
9376	79636042 (7025, 7026)			UNCLASSIFIED	1023

9377	65702705 (7027, 7028)			UNCLASSIFIED	1038
9378	54536815 (7037, 7038)			UNCLASSIFIED	1037
9379	71204317 (7039, 7040)			UNCLASSIFIED	1021, 1050
9380	78404994 (7051, 7052)			UNCLASSIFIED	1016, 1024
9381	71846635 (7073, 7074)			UNCLASSIFIED	1024
9382	86467178 (7079, 7080)			UNCLASSIFIED	1029
9383	86473907 (7083, 7084)			UNCLASSIFIED	1029
9384	71037505 (7091, 7092)			UNCLASSIFIED	1034, 1054
9385	70618824 (7101, 7102)			UNCLASSIFIED	1036
9386	54526537 (7107, 7108)			UNCLASSIFIED	1036
9387	95005590 (7111, 7112)			UNCLASSIFIED	1034
9388	11384272 (7113, 7114)			UNCLASSIFIED	1016
9389	94328529 (7117, 7118)			UNCLASSIFIED	1054
9390	29445365 (7119, 7120)			UNCLASSIFIED	1038
9391	80051084 (7123, 7124)			UNCLASSIFIED	1004
9392	11777094 (7125, 7126)			UNCLASSIFIED	1038
9393	78297024 (7133, 7134)			UNCLASSIFIED	1026
9394	79582823 (7135, 7136)			UNCLASSIFIED	1027

9395	54701874 (7159, 7160)			UNCLASSIFIED	1019
9396	91213991 (7161, 7162)			UNCLASSIFIED	1054
9397	49263153 (7165, 7166)			UNCLASSIFIED	1040
9398	35129859 (7173, 7174)			UNCLASSIFIED	1012
9399	80059599 (7177, 7178)			UNCLASSIFIED	1010
9400	35994013 (7179, 7180)			UNCLASSIFIED	1024
9401	43336687 (7183, 7184)			UNCLASSIFIED	1000
9402	65700668 (7197, 7198)			UNCLASSIFIED	1038
9403	30174748 (7215, 7216)			UNCLASSIFIED	1030
9404	85801391 (7221, 7222)			UNCLASSIFIED	1038
9405	78979641 (7233, 7234)			UNCLASSIFIED	1016, 1050
9406	13761567 (7235, 7236)			UNCLASSIFIED	1016
9407	70944560 (7239, 7240)			UNCLASSIFIED	1024
9408	10858857 (7255, 7256)			UNCLASSIFIED	1031
9409	91237478 (7269, 7270)			UNCLASSIFIED	1000, 1010
9410	8787371 (7275, 7276)			UNCLASSIFIED	1022
9411	29456966 (7285, 7286)			UNCLASSIFIED	1026
9412	24144352 (7291, 7292)			UNCLASSIFIED	1003

9413	56776332 (7297, 7298)			UNCLASSIFIED	1042
9414	71034349 (7305, 7306)			UNCLASSIFIED	1026
9415	66143035 (7307, 7308)			UNCLASSIFIED	1010
9416	33200750 (7311, 7312)			UNCLASSIFIED	1026
9417	66163918 (7331, 7332)			UNCLASSIFIED	1024
9418	34407275 (7341, 7342)			UNCLASSIFIED	1038
9419	33099596 (7347, 7348)			UNCLASSIFIED	1026
9420	79485063 (7349, 7350)			UNCLASSIFIED	1016
9421	94232484 (7357, 7358)			UNCLASSIFIED	1054
9422	39523113 (7361, 7362)			UNCLASSIFIED	1022
9423	66373527 (7363, 7364)			UNCLASSIFIED	1054
9424	49623727 (7367, 7368)			UNCLASSIFIED	1036
9425	11812483 (7389, 7390)			UNCLASSIFIED	1004
9426	47656774 (7411, 7412)			UNCLASSIFIED	1054
9427	11398745 (7421, 7422)			UNCLASSIFIED	1006
9428	79244505 (7425, 7426)			UNCLASSIFIED	1024, 1038
9429	79557077 (7435, 7436)			UNCLASSIFIED	1025
9430	65985516 (7443, 7444)			UNCLASSIFIED	1054

9431	33764770 (7445, 7446)			UNCLASSIFIED	1026
9432	16841837 (7465, 7466)			UNCLASSIFIED	1044
9433	79312124 (7469, 7470)			UNCLASSIFIED	1004
9434	87456054 (7475, 7476)			UNCLASSIFIED	1025
9435	66173591 (7498, 7500)			UNCLASSIFIED	1016
9436	17296223 (7503, 7504)			UNCLASSIFIED	1003
9437	21635874 (7507, 7508)			UNCLASSIFIED	1044
9438	79493084 (7511, 7512)			UNCLASSIFIED	1006, 1044
9439	57057454 (7513, 7514)			UNCLASSIFIED	1051
9440	34754674 (7523, 7524)			UNCLASSIFIED	1024
9441	78386715 (7525, 7526)			UNCLASSIFIED	1023, 1026
9442	39523739 (7527, 7528)			UNCLASSIFIED	1004
9443	21431341 (7531, 7532)			UNCLASSIFIED	1004
9444	11102975 (7535, 7536)			UNCLASSIFIED	1022
9445	33763086 (7537, 7538)			UNCLASSIFIED	1026
9446	33047831 (7539, 7540)			UNCLASSIFIED	1026
9447	66256508 (7543, 7544)			UNCLASSIFIED	1016
9448	47652855 (7555, 7556)			UNCLASSIFIED	1029

9449	21430323 (7579, 7580)			UNCLASSIFIED	1010
9450	86688638 (7599, 7600)			UNCLASSIFIED	1029
9451	20294573 (7603, 7604)			UNCLASSIFIED	1034
9452	86477452 (7619, 7620)			UNCLASSIFIED	1025, 1029
9453	78396194 (7627, 7628)			UNCLASSIFIED	1021, 1024, 1030
9454	17887616 (7637, 7638)			UNCLASSIFIED	1024
9455	6324745 (7641, 7642)			UNCLASSIFIED	1058
9456	28860851 (7645, 7646)			UNCLASSIFIED	1003
9457	25259181 (7649, 7650)			UNCLASSIFIED	1003
9458	66317462 (7667, 7668)			UNCLASSIFIED	1038
9459	43154549 (7671, 7672)			UNCLASSIFIED	1001
9460	79637077 (7679, 7680)			UNCLASSIFIED	1038
9461	79552099 (7703, 7704)			UNCLASSIFIED	1025
9462	65700443 (7705, 7706)			UNCLASSIFIED	1038
9463	79560386 (7707, 7708)			UNCLASSIFIED	1038
9464	78963606 (7709, 7710)			UNCLASSIFIED	1038, 1054
9465	10997264 (7759, 7760)			UNCLASSIFIED	1024
9466	70761103 (7765, 7766)			UNCLASSIFIED	1023

9467	79639676 (7775, 7776)			UNCLASSIFIED	1019, 1023, 1025, 1030
9468	9387387 (7781, 7782)			UNCLASSIFIED	1003
9469	80054978 (7789, 7790)			UNCLASSIFIED	1004
9470	65462323 (7791, 7792)			UNCLASSIFIED	1054
9471	36902755 (7793, 7794)			UNCLASSIFIED	1026
9472	38399024 (7803, 7804)			UNCLASSIFIED	1026
9473	95295646 (7807, 7808)			UNCLASSIFIED	1025
9474	94749190 (7821, 7822)			UNCLASSIFIED	1044
9475	70620029 (7825, 7826)			UNCLASSIFIED	1036
9476	79627253 (7827, 7828)			UNCLASSIFIED	1038
9477	65445326 (7829, 7830)			UNCLASSIFIED	1016
9478	66173775 (7833, 7834)			UNCLASSIFIED	1016
9479	86380037 (7857, 7858)			UNCLASSIFIED	1053
9480	94854079 (7865, 7866)			UNCLASSIFIED	1010
9481	78505846 (7867, 7868)			UNCLASSIFIED	1016
9482	34115090 (7871, 7872)			UNCLASSIFIED	1000
9483	46891805 (7881, 7882)			UNCLASSIFIED	1019
9484	21419100 (7883, 7884)			UNCLASSIFIED	1010

9485	78977278 (7891, 7892)			UNCLASSIFIED	1016
9486	65695325 (7903, 7904)			UNCLASSIFIED	1026, 1038, 1044
9487	33204895 (7907, 7908)			UNCLASSIFIED	1026
9488	79635453 (7913, 7914)			UNCLASSIFIED	1038
9489	65981609 (7915, 7916)			UNCLASSIFIED	1054
9490	71203122 (7917, 7918)			UNCLASSIFIED	1030, 1038
9491	25143236 (7921, 7922)			UNCLASSIFIED	1008
9492	27842126 (7925, 7926)			UNCLASSIFIED	1006
9493	80067429 (7929, 7930)			UNCLASSIFIED	1034
9494	80210773 (7933, 7934)			UNCLASSIFIED	1000, 1006, 1024, 1026, 1034
9495	27975661 (7943, 7944)			UNCLASSIFIED	1022
9496	78945294 (7947, 7948)			UNCLASSIFIED	1030
9497	10220696 (7953, 7954)			UNCLASSIFIED	1039
9498	29910808 (7955, 7956)			UNCLASSIFIED	1030
9499	79212700 (7967, 7968)			UNCLASSIFIED	1017, 1023, 1054
9500	25259185 (7971, 7972)			UNCLASSIFIED	1026
9501	85811409 (7993, 7994)			UNCLASSIFIED	1019
9502	47656894 (7997, 7998)			UNCLASSIFIED	1029

9503	35997383 (7999, 8000)			UNCLASSIFIED	1029
9504	16759984 (8005, 8006)			UNCLASSIFIED	1023
9505	37801394 (8015, 8016)			UNCLASSIFIED	1012
9506	21629804 (8023, 8024)			UNCLASSIFIED	1044
9507	79389814 (8033, 8034)			UNCLASSIFIED	1023
9508	86477457 (8037, 8038)			UNCLASSIFIED	1029
9509	78742212 (8039, 8040)			UNCLASSIFIED	1044
9510	79458186 (8047, 8048)			UNCLASSIFIED	1031, 1051
9511	80048406 (8049, 8050)			UNCLASSIFIED	1024, 1025
9512	78746295 (8055, 8056)			UNCLASSIFIED	1017, 1030
9513	65497629 (8063, 8064)			UNCLASSIFIED	1016
9514	19638228 (8065, 8066)			UNCLASSIFIED	1054
9515	79574277 (8067, 8068)			UNCLASSIFIED	1004, 1006, 1031
9516	65685880 (8075, 8076)			UNCLASSIFIED	1024
9517	28833779 (8087, 8088)			UNCLASSIFIED	1022
9518	79633344 (8089, 8090)			UNCLASSIFIED	1021, 1038
9519	20729558 (8091, 8092)			UNCLASSIFIED	1004
9520	80188399 (8095, 8096)			UNCLASSIFIED	1058

9521	11076189 (8099, 8100)			UNCLASSIFIED	1022
9522	79817382 (8111, 8112)			UNCLASSIFIED	1017
9523	80503816 (8113, 8114)			UNCLASSIFIED	1004, 1012, 1017
9524	78769303 (8119, 8120)			UNCLASSIFIED	1003, 1017
9525	32358605 (8123, 8124)			UNCLASSIFIED	1054
9526	78500167 (8133, 8134)			UNCLASSIFIED	1038
9527	71220972 (8153, 8154)			UNCLASSIFIED	1000
9528	65466386 (8155, 8156)			UNCLASSIFIED	1022, 1054
9529	33183219 (8167, 8168)			UNCLASSIFIED	1026
9530	9857184 (8171, 8172)			UNCLASSIFIED	1008
9531	13508904 (8175, 8176)			UNCLASSIFIED	1034
9532	78483746 (8177, 8178)			UNCLASSIFIED	1026
9533	86466029 (8187, 8188)			UNCLASSIFIED	1029
9534	36997745 (8195, 8196)			UNCLASSIFIED	1001
9535	78961540 (8209, 8210)			UNCLASSIFIED	1054
9536	21415299 (8213, 8214)			UNCLASSIFIED	1034
9537	82120206 (8217, 8218)			UNCLASSIFIED	1016, 1031
9538	78762612 (8223, 8224)			UNCLASSIFIED	1044

9539	78529072 (8227, 8228)			UNCLASSIFIED	1026
9540	28458757 (8235, 8236)			UNCLASSIFIED	1030
9541	65644212 (8237, 8238)			UNCLASSIFIED	1041
9542	78946817 (8253, 8254)			UNCLASSIFIED	1003, 1009
9543	55780076 (8267, 8268)			UNCLASSIFIED	1008
9544	47657004 (8269, 8270)			UNCLASSIFIED	1029
9545	9244967 (8271, 8272)			UNCLASSIFIED	1017
9546	78302743 (8279, 8280)			UNCLASSIFIED	1026
9547	10108819 (8287, 8288)			UNCLASSIFIED	1044
9548	10154780 (8307, 8308)			UNCLASSIFIED	1006
9549	78678841 (8319, 8320)			UNCLASSIFIED	1003
9550	20606803 (8335, 8336)			UNCLASSIFIED	1004
9551	66064097 (8345, 8346)			UNCLASSIFIED	1008
9552	80023353 (8377, 8378)			UNCLASSIFIED	1022
9553	34408568 (8379, 8380)			UNCLASSIFIED	1026
9554	79963951 (8405, 8406)			UNCLASSIFIED	1003, 1006, 1021, 1025
9555	20290083 (8413, 8414)			UNCLASSIFIED	1022
9556	16410019 (8425, 8426)			UNCLASSIFIED	1054

9557	78729835 (8437, 8438)			UNCLASSIFIED	1022, 1023, 1026
9558	78503290 (8445, 8446)			UNCLASSIFIED	1026
9559	55180170 (8455, 8456)			UNCLASSIFIED	1019
9560	57458269 (8463, 8464)			UNCLASSIFIED	1054
9561	34269759 (8465, 8466)			UNCLASSIFIED	1026
9562	27990468 (8487, 8488)			UNCLASSIFIED	1025
9563	27968206 (8493, 8494)			UNCLASSIFIED	1006
9564	20733682 (8499, 8500)			UNCLASSIFIED	1022
9565	78459336 (8501, 8502)			UNCLASSIFIED	1026
9566	9863861 (8503, 8504)			UNCLASSIFIED	1008
9567	79869898 (8507, 8508)			UNCLASSIFIED	1017
9568	8748466 (8515, 8516)			UNCLASSIFIED	1010
9569	20606807 (8523, 8524)			UNCLASSIFIED	1024
9570	38281598 (8531, 8532)			UNCLASSIFIED	1016
9571	19508297 (8537, 8538)			UNCLASSIFIED	1034
9572	33694193 (8543, 8544)			UNCLASSIFIED	1026
9573	33184569 (8555, 8556)			UNCLASSIFIED	1026
9574	57441293 (8557, 8558)			UNCLASSIFIED	1058

9575	29224499 (8571, 8572)			UNCLASSIFIED	1030
9576	23282274 (8575, 8576)			UNCLASSIFIED	1003, 1022
9577	79638143 (8577, 8578)			UNCLASSIFIED	1044
9578	56068549 (8579, 8580)			UNCLASSIFIED	1021
9579	20270200 (8585, 8586)			UNCLASSIFIED	1010
9580	79170469 (8599, 8600)			UNCLASSIFIED	1024, 1039
9581	79561016 (8601, 8602)			UNCLASSIFIED	1044
9582	78485310 (8607, 8608)			UNCLASSIFIED	1026
9583	71228724 (8617, 8618)			UNCLASSIFIED	1000
9584	20466350 (8629, 8630)			UNCLASSIFIED	1010
9585	66437329 (8631, 8632)			UNCLASSIFIED	1024
9586	49136629 (8637, 8638)			UNCLASSIFIED	1036, 1054
9587	17658173 (8639, 8640)			UNCLASSIFIED	1025
9588	78793149 (8643, 8644)			UNCLASSIFIED	1044
9589	55663198 (8645, 8646)			UNCLASSIFIED	1016
9590	8755542 (8647, 8648)			UNCLASSIFIED	1004
9591	77522297 (8659, 8660)			UNCLASSIFIED	1013
9592	95090596 (8663, 8664)			UNCLASSIFIED	1044

9593	82310070 (8671, 8672)			UNCLASSIFIED	1001
9594	66644928 (8685, 8686)			UNCLASSIFIED	1016
9595	54531892 (8687, 8688)			UNCLASSIFIED	1041
9596	66672118 (8689, 8690)			UNCLASSIFIED	1016
9597	78256716 (8691, 8692)			UNCLASSIFIED	1026
9598	20630346 (8697, 8698)			UNCLASSIFIED	1004
9599	79115216 (8701, 8702)			UNCLASSIFIED	1024
9600	57524197 (8705, 8706)			UNCLASSIFIED	1006, 1010, 1024, 1025
9601	71088771 (8727, 8728)			UNCLASSIFIED	1053
9602	78386692 (8729, 8730)			UNCLASSIFIED	1026
9603	79827816 (8731, 8732)			UNCLASSIFIED	1003, 1024
9604	56658960 (8745, 8746)			UNCLASSIFIED	1008
9605	78146449 (8751, 8752)			UNCLASSIFIED	1026
9606	80025710 (8755, 8756)			UNCLASSIFIED	1022, 1034, 1044
9607	91254738 (8765, 8766)			UNCLASSIFIED	1010, 1038, 1040
9608	42563300 (8771, 8772)			UNCLASSIFIED	1037
9609	23300350 (8775, 8776)			UNCLASSIFIED	1009
9610	23288792 (8781, 8782)			UNCLASSIFIED	1001, 1023

9611	54527942 (8787, 8788)			UNCLASSIFIED	1041
9612	27797150 (8795, 8796)			UNCLASSIFIED	1022
9613	79178083 (8797, 8798)			UNCLASSIFIED	1003, 1008, 1017, 1024
9614	27365748 (8805, 8806)			UNCLASSIFIED	1017
9615	39439118 (8809, 8810)			UNCLASSIFIED	1054
9616	78736064 (8817, 8818)			UNCLASSIFIED	1003, 1023, 1039, 1044
9617	65888459 (8837, 8838)			UNCLASSIFIED	1054
9618	10330714 (8843, 8844)			UNCLASSIFIED	1025
9619	33108424 (8847, 8848)			UNCLASSIFIED	1000
9620	78360965 (8857, 8858)			UNCLASSIFIED	1008
9621	79612364 (8879, 8880)			UNCLASSIFIED	1039
9622	29691237 (8881, 8882)			UNCLASSIFIED	1024
9623	52212590 (8885, 8886)			UNCLASSIFIED	1040
9624	57413514 (8889, 8890)			UNCLASSIFIED	1000, 1058
9625	77932535 (8895, 8896)			UNCLASSIFIED	1036
9626	7425974 (8905, 8906)			UNCLASSIFIED	1058
9627	23303585 (8913, 8914)			UNCLASSIFIED	1023
9628	77952824 (8917, 8918)			UNCLASSIFIED	1036

9629	80422615 (8925, 8926)			UNCLASSIFIED	1024, 1025, 1038, 1054
9630	10211489 (8935, 8936)			UNCLASSIFIED	1004
9631	79189901 (8947, 8948)			UNCLASSIFIED	1024
9632	16329719 (8949, 8950)			UNCLASSIFIED	1004
9633	27824678 (8955, 8956)			UNCLASSIFIED	1010
9634	29448376 (8965, 8966)			UNCLASSIFIED	1022
9635	20417410 (8969, 8970)			UNCLASSIFIED	1024
9636	86667079 (8979, 8980)			UNCLASSIFIED	1009, 1029
9637	33700910 (8981, 8982)			UNCLASSIFIED	1026
9638	78895259 (8995, 8996)			UNCLASSIFIED	1003
9639	71781964 (8997, 8998)			UNCLASSIFIED	1044
9640	20449239 (9001, 9002)			UNCLASSIFIED	1010
9641	65849023 (9009, 9010)			UNCLASSIFIED	1038
9642	28810462 (9025, 9026)			UNCLASSIFIED	1022
9643	78401861 (9027, 9028)			UNCLASSIFIED	1016, 1024, 1054
9644	78980871 (9029, 9030)			UNCLASSIFIED	1016
9645	20292971 (9031, 9032)			UNCLASSIFIED	1034
9646	9285408 (9035, 9036)			UNCLASSIFIED	1003

9647	46888531 (9047, 9048)			UNCLASSIFIED	1037
9648	91229654 (9049, 9050)			UNCLASSIFIED	1010
9649	30244035 (9055, 9056)			UNCLASSIFIED	1026
9650	32275665 (9083, 9084)			UNCLASSIFIED	1029
9651	39444092 (9087, 9088)			UNCLASSIFIED	1022
9652	79633668 (9097, 9098)			UNCLASSIFIED	1038
9653	79254311 (9103, 9104)			UNCLASSIFIED	1023
9654	27849916 (9109, 9110)			UNCLASSIFIED	1022
9655	79557085 (9111, 9112)			UNCLASSIFIED	1025
9656	57460711 (9115, 9116)			UNCLASSIFIED	1011
9657	27959991 (9127, 9128)			UNCLASSIFIED	1022
9658	80062402 (9137, 9138)			UNCLASSIFIED	1010
9659	10212686 (9139, 9140)			UNCLASSIFIED	1025
9660	33184986 (9143, 9144)			UNCLASSIFIED	1026
9661	30655242 (9165, 9166)			UNCLASSIFIED	1022
9662	27348915 (9169, 9170)			UNCLASSIFIED	1026
9663	16278600 (9171, 9172)			UNCLASSIFIED	1006
9664	20459498 (9175, 9176)			UNCLASSIFIED	1010

9665	78890098 (9179, 9180)			UNCLASSIFIED	1003
9666	46874324 (9185, 9186)			UNCLASSIFIED	1037
9667	16345938 (9193, 9194)			UNCLASSIFIED	1034
9668	78384654 (9207, 9208)			UNCLASSIFIED	1017
9669	29210581 (9211, 9212)			UNCLASSIFIED	1017
9670	28486710 (9215, 9216)			UNCLASSIFIED	1026
9671	79562426 (9227, 9228)			UNCLASSIFIED	1038
9672	65479717 (9229, 9230)			UNCLASSIFIED	1054
9673	79652728 (9239, 9240)			UNCLASSIFIED	1001, 1006, 1022, 1023, 1024, 1026
9674	25146776 (9263, 9264)			UNCLASSIFIED	1003
9675	32311056 (9295, 9296)			UNCLASSIFIED	1024
9676	25305744 (9301, 9302)			UNCLASSIFIED	1003
9677	20595869 (9303, 9304)			UNCLASSIFIED	1004
9678	80069364 (9307, 9308)			UNCLASSIFIED	1022, 1040, 1046
9679	10315991 (9313, 9314)			UNCLASSIFIED	1003
9680	32309370 (9321, 9322)			UNCLASSIFIED	1029
9681	66176330 (9329, 9330)			UNCLASSIFIED	1016
9682	54994459 (9339, 9340)			UNCLASSIFIED	1029

9683	78254513 (9355, 9356)			UNCLASSIFIED	1026
9684	79584629 (9357, 9358)			UNCLASSIFIED	1024
9685	78790882 (9359, 9360)			UNCLASSIFIED	1008
9686	32684069 (9367, 9368)			UNCLASSIFIED	1026
9687	80186595 (9371, 9372)			UNCLASSIFIED	1054, 1058
9688	78982140 (9375, 9376)			UNCLASSIFIED	1030
9689	14401563 (9379, 9380)			UNCLASSIFIED	1053
9690	36518176 (9393, 9394)			UNCLASSIFIED	1034
9691	78992752 (9395, 9396)			UNCLASSIFIED	1044
9692	20435084 (9407, 9408)			UNCLASSIFIED	1022
9693	79853312 (9413, 9414)			UNCLASSIFIED	1017
9694	10103677 (9443, 9444)			UNCLASSIFIED	1039
9695	81226684 (9449, 9450)			UNCLASSIFIED	1003, 1023, 1044, 1054
9696	20709887 (9455, 9456)			UNCLASSIFIED	1022
9697	54859987 (9459, 9460)			UNCLASSIFIED	1017, 1019, 1041
9698	13513092 (9473, 9474)			UNCLASSIFIED	1024
9699	78796794 (9475, 9476)			UNCLASSIFIED	1026
9700	86684849 (9477, 9478)			UNCLASSIFIED	1029

9701	80083946 (9489, 9490)			UNCLASSIFIED	1001
9702	86380050 (9495, 9496)			UNCLASSIFIED	1053
9703	80062406 (9501, 9502)			UNCLASSIFIED	1010, 1022, 1027
9704	79261079 (9503, 9504)			UNCLASSIFIED	1024
9705	30269847 (9509, 9510)			UNCLASSIFIED	1026
9706	86675948 (9521, 9522)			UNCLASSIFIED	1007
9707	80033111 (9523, 9524)			UNCLASSIFIED	1053
9708	79840407 (9531, 9532)			UNCLASSIFIED	1001, 1008, 1044, 1049, 1054
9709	77727488 (9533, 9534)			UNCLASSIFIED	1016
9710	78252606 (9537, 9538)			UNCLASSIFIED	1012
9711	66049476 (9549, 9550)			UNCLASSIFIED	1007
9712	33185021 (9555, 9556)			UNCLASSIFIED	1026
9713	34624669 (9557, 9558)			UNCLASSIFIED	1029
9714	85805692 (9565, 9566)			UNCLASSIFIED	1019, 1036
9715	35110330 (9571, 9572)			UNCLASSIFIED	1029
9716	30409127 (9577, 9578)			UNCLASSIFIED	1026
9717	80503029 (9579, 9580)			UNCLASSIFIED	1012, 1054
9718	19890416 (9589, 9590)			UNCLASSIFIED	1054

9719	28398479 (9601, 9602)			UNCLASSIFIED	1008
9720	65645704 (9603, 9604)			UNCLASSIFIED	1041
9721	50843727 (9615, 9616)			UNCLASSIFIED	1036
9722	79255725 (9617, 9618)			UNCLASSIFIED	1025
9723	29519358 (9621, 9622)			UNCLASSIFIED	1006
9724	46764504 (9627, 9628)			UNCLASSIFIED	1054
9725	18166077 (9629, 9630)			UNCLASSIFIED	1038
9726	65468180 (9631, 9632)			UNCLASSIFIED	1054
9727	56516549 (9633, 9634)			UNCLASSIFIED	1019
9728	78979791 (9637, 9638)			UNCLASSIFIED	1016
9729	29444352 (9651, 9652)			UNCLASSIFIED	1026
9730	29884721 (9655, 9656)			UNCLASSIFIED	1024
9731	79604352 (9657, 9658)			UNCLASSIFIED	1003
9732	25261403 (9669, 9670)			UNCLASSIFIED	1008
9733	70961131 (9671, 9672)			UNCLASSIFIED	1000, 1019
9734	78507898 (9679, 9680)			UNCLASSIFIED	1009
9735	27362480 (9685, 9686)			UNCLASSIFIED	1017
9736	66428042 (9691, 9692)			UNCLASSIFIED	1022

9737	29916701 (9701, 9702)			UNCLASSIFIED	1030
9738	29213107 (9713, 9714)			UNCLASSIFIED	1017
9739	66049961 (9715, 9716)			UNCLASSIFIED	1054
9740	78379813 (9717, 9718)			UNCLASSIFIED	1026
9741	16410363 (9727, 9728)			UNCLASSIFIED	1003
9742	10063185 (9731, 9732)			UNCLASSIFIED	1017
9743	79256651 (9739, 9740)			UNCLASSIFIED	1029
9744	80239396 (9749, 9750)			UNCLASSIFIED	1004, 1006, 1010, 1023, 1024
9745	54700304 (9753, 9754)			UNCLASSIFIED	1022
9746	80101757 (9765, 9766)			UNCLASSIFIED	1022, 1024
9747	78752705 (9769, 9770)			UNCLASSIFIED	1023, 1039
9748	65683216 (9775, 9776)			UNCLASSIFIED	1024
9749	56156500 (9777, 9778)			UNCLASSIFIED	1009, 1023
9750	20084281 (9783, 9784)			UNCLASSIFIED	1034
9751	78254181 (9789, 9790)			UNCLASSIFIED	1026
9752	35061224 (9801, 9802)			UNCLASSIFIED	1054
9753	65701172 (9803, 9804)			UNCLASSIFIED	1038
9754	34765342 (9811, 9812)			UNCLASSIFIED	1050

9755	80200459 (9819, 9820)			UNCLASSIFIED	1022
9756	29681435 (9827, 9828)			UNCLASSIFIED	1038
9757	80503554 (9829, 9830)			UNCLASSIFIED	1004, 1006, 1010, 1012, 1017, 1022, 1024, 1038, 1044
9758	29247142 (9845, 9846)			UNCLASSIFIED	1026
9759	78500310 (9861, 9862)			UNCLASSIFIED	1018
9760	20631931 (9865, 9866)			UNCLASSIFIED	1024
9761	32126051 (9867, 9868)			UNCLASSIFIED	1024
9762	28002248 (9875, 9876)			UNCLASSIFIED	1054
9763	79234049 (9883, 9884)			UNCLASSIFIED	1001
9764	78254893 (9891, 9892)			UNCLASSIFIED	1026
9765	8242670 (9895, 9896)			UNCLASSIFIED	1034
9766	20284095 (9901, 9902)			UNCLASSIFIED	1022
9767	78399092 (9905, 9906)			UNCLASSIFIED	1016
9768	17898139 (9907, 9908)			UNCLASSIFIED	1044
9769	85812169 (9933, 9934)			UNCLASSIFIED	1024
9770	30789290 (9935, 9936)			UNCLASSIFIED	1026
9771	14998191 (9941, 9942)			UNCLASSIFIED	1024

9772	35893386 (9945, 9946)			UNCLASSIFIED	1053
9773	87460159 (9947, 9948)			UNCLASSIFIED	1048
9774	85531528 (9949, 9950)			UNCLASSIFIED	1038
9775	25330495 (9965, 9966)			UNCLASSIFIED	1003
9776	30673193 (9971, 9972)			UNCLASSIFIED	1008
9777	78911398 (9977, 9978)			UNCLASSIFIED	1044
9778	79631762 (9981, 9982)			UNCLASSIFIED	1024, 1044
9779	38898732 (9997, 9998)			UNCLASSIFIED	1022
9780	78049724 (10009, 10010)			UNCLASSIFIED	1026
9781	87465866 (10023, 10024)			UNCLASSIFIED	1018
9782	94846244 (10025, 10026)			UNCLASSIFIED	1044
9783	50876837 (10037, 10038)			UNCLASSIFIED	1036
9784	55401531 (10047, 10048)			UNCLASSIFIED	1019
9785	77921355 (10055, 10056)			UNCLASSIFIED	1041
9786	71046211 (10061, 10062)			UNCLASSIFIED	1016, 1024
9787	86666685 (10087, 10088)			UNCLASSIFIED	1004, 1026, 1029
9788	91235589 (10099, 10100)			UNCLASSIFIED	1010
9789	46847289 (10111, 10112)			UNCLASSIFIED	1037

9790	20728293 (10119, 10120)			UNCLASSIFIED	1006
9791	79634211 (10121, 10122)			UNCLASSIFIED	1038
9792	28326379 (10135, 10136)			UNCLASSIFIED	1038
9793	78074770 (10149, 10150)			UNCLASSIFIED	1024
9794	79760476 (10151, 10152)			UNCLASSIFIED	1008
9795	80027628 (10155, 10156)			UNCLASSIFIED	1004, 1031
9796	32682120 (10159, 10160)			UNCLASSIFIED	1026
9797	78254642 (10161, 10162)			UNCLASSIFIED	1026
9798	10156843 (10169, 10170)			UNCLASSIFIED	1025
9799	85512920 (10175, 10176)			UNCLASSIFIED	1053
9800	65893410 (10177, 10178)			UNCLASSIFIED	1054
9801	79612034 (10181, 10182)			UNCLASSIFIED	1006
9802	65700052 (10197, 10198)			UNCLASSIFIED	1038
9803	79559859 (10199, 10200)			UNCLASSIFIED	1024, 1038
9804	87447932 (10201, 10202)			UNCLASSIFIED	1018
9805	78910592 (10203, 10204)			UNCLASSIFIED	1039
9806	65452282 (10207, 10208)			UNCLASSIFIED	1016
9807	20754894 (10221, 10222)			UNCLASSIFIED	1053

9808	78983211 (10223, 10224)			UNCLASSIFIED	1054
9809	78369434 (10235, 10236)			UNCLASSIFIED	1017
9810	79568849 (10243, 10244)			UNCLASSIFIED	1008
9811	46688011 (10245, 10246)			UNCLASSIFIED	1022
9812	78921245 (10255, 10256)			UNCLASSIFIED	1017
9813	80071706 (10259, 10260)			UNCLASSIFIED	1003, 1004, 1006, 1022
9814	29442886 (10265, 10266)			UNCLASSIFIED	1026
9815	78526436 (10267, 10268)			UNCLASSIFIED	1009
9816	30253868 (10269, 10270)			UNCLASSIFIED	1026
9817	19848552 (10271, 10272)			UNCLASSIFIED	1034
9818	33187811 (10273, 10274)			UNCLASSIFIED	1026
9819	66762861 (10287, 10288)			UNCLASSIFIED	1024
9820	20447626 (10311, 10312)			UNCLASSIFIED	1010
9821	38399132 (10313, 10314)			UNCLASSIFIED	1026
9822	39550350 (10327, 10328)			UNCLASSIFIED	1038
9823	38069707 (10333, 10334)			UNCLASSIFIED	1016
9824	29508303 (10335, 10336)			UNCLASSIFIED	1024
9825	78771536 (10345, 10346)			UNCLASSIFIED	1008

9826	11086180 (10347, 10348)			UNCLASSIFIED	1034
9827	79831686 (10349, 10350)			UNCLASSIFIED	1008
9828	79573792 (10351, 10352)			UNCLASSIFIED	1001
9829	35050262 (10353, 10354)			UNCLASSIFIED	1030
9830	34412301 (10367, 10368)			UNCLASSIFIED	1050
9831	52496092 (10375, 10376)			UNCLASSIFIED	1036
9832	33706090 (10391, 10392)			UNCLASSIFIED	1050
9833	20817572 (10393, 10394)			UNCLASSIFIED	1004
9834	16292974 (10399, 10400)			UNCLASSIFIED	1054
9835	78780816 (10407, 10408)			UNCLASSIFIED	1026
9836	30376078 (10419, 10420)			UNCLASSIFIED	1026
9837	55177125 (10425, 10426)			UNCLASSIFIED	1019
9838	86473268 (10429, 10430)			UNCLASSIFIED	1029
9839	78747674 (10437, 10438)			UNCLASSIFIED	1026
9840	10886616 (10447, 10448)			UNCLASSIFIED	1022
9841	70441655 (10461, 10462)			UNCLASSIFIED	1019, 1021, 1041
9842	78363896 (10465, 10466)			UNCLASSIFIED	1008
9843	85811423 (10483, 10484)			UNCLASSIFIED	1019

9844	78385850 (10485, 10486)			UNCLASSIFIED	1044
9845	78460294 (10493, 10494)			UNCLASSIFIED	1003
9846	19848700 (10499, 10500)			UNCLASSIFIED	1022
9847	20431007 (10501, 10502)			UNCLASSIFIED	1004, 1023
9848	24140029 (10503, 10504)			UNCLASSIFIED	1003, 1039
9849	66138814 (10517, 10518)			UNCLASSIFIED	1004
9850	29466863 (10523, 10524)			UNCLASSIFIED	1038
9851	55186371 (10535, 10536)			UNCLASSIFIED	1019
9852	65685013 (10537, 10538)			UNCLASSIFIED	1046
9853	20287405 (10541, 10542)			UNCLASSIFIED	1022
9854	25242190 (10545, 10546)			UNCLASSIFIED	1008
9855	35993579 (10549, 10550)			UNCLASSIFIED	1024
9856	27978395 (10559, 10560)			UNCLASSIFIED	1022
9857	55181579 (10561, 10562)			UNCLASSIFIED	1019
9858	86666169 (10563, 10564)			UNCLASSIFIED	1007
9859	78256728 (10565, 10566)			UNCLASSIFIED	1026
9860	91215973 (10573, 10574)			UNCLASSIFIED	1053
9861	9285106 (10577, 10578)			UNCLASSIFIED	1008

9862	94670425 (10581, 10582)			UNCLASSIFIED	1024
9863	78245903 (10593, 10594)			UNCLASSIFIED	1022, 1026
9864	91232226 (10601, 10602)			UNCLASSIFIED	1010
9865	80080496 (10603, 10604)			UNCLASSIFIED	1034
9866	10095590 (10607, 10608)			UNCLASSIFIED	1044
9867	35916807 (10615, 10616)			UNCLASSIFIED	1022
9868	54693746 (10619, 10620)			UNCLASSIFIED	1041
9869	32445733 (10631, 10632)			UNCLASSIFIED	1001
9870	27804573 (10635, 10636)			UNCLASSIFIED	1010
9871	9285092 (10651, 10652)			UNCLASSIFIED	1003
9872	78307296 (10655, 10656)			UNCLASSIFIED	1026
9873	78788922 (10661, 10662)			UNCLASSIFIED	1009, 1010, 1054
9874	80034661 (10671, 10672)			UNCLASSIFIED	1053
9875	33694203 (10673, 10674)			UNCLASSIFIED	1026
9876	78679394 (10677, 10678)			UNCLASSIFIED	1017
9877	87449678 (10679, 10680)			UNCLASSIFIED	1025
9878	38444969 (10713, 10714)			UNCLASSIFIED	1026
9879	55181729 (10727, 10728)			UNCLASSIFIED	1019

9880	33187128 (10733, 10734)			UNCLASSIFIED	1026
9881	1499206 (10741, 10742)			UNCLASSIFIED	1024
9882	29354857 (10743, 10744)			UNCLASSIFIED	1009
9883	80042037 (10749, 10750)			UNCLASSIFIED	1024
9884	80069083 (10753, 10754)			UNCLASSIFIED	1010, 1024
9885	30179178 (10755, 10756)			UNCLASSIFIED	1029
9886	78680342 (10771, 10772)			UNCLASSIFIED	1017, 1033, 1054, 1055
9887	80058744 (10785, 10786)			UNCLASSIFIED	1010, 1022
9888	79599699 (10787, 10788)			UNCLASSIFIED	1003
9889	25252692 (10803, 10804)			UNCLASSIFIED	1008
9890	79314952 (10805, 10806)			UNCLASSIFIED	1013, 1019, 1022, 1038
9891	85539439 (10811, 10812)			UNCLASSIFIED	1054
9892	17927790 (10823, 10824)			UNCLASSIFIED	1038
9893	79871220 (10857, 10858)			UNCLASSIFIED	1017
9894	71749048 (10859, 10860)			UNCLASSIFIED	1003, 1044
9895	13527826 (10861, 10862)			UNCLASSIFIED	1001
9896	27921494 (10865, 10866)			UNCLASSIFIED	1022, 1039
9897	78293455 (10867, 10868)			UNCLASSIFIED	1026, 1030

9898	71230414 (10881, 10882)			UNCLASSIFIED	1016, 1050
9899	30407335 (10903, 10904)			UNCLASSIFIED	1026
9900	23302664 (10905, 10906)			UNCLASSIFIED	1026
9901	80239084 (10913, 10914)			UNCLASSIFIED	1010, 1017
9902	37019834 (10927, 10928)			UNCLASSIFIED	1025
9903	33266443 (10947, 10948)			UNCLASSIFIED	1029
9904	94256063 (10951, 10952)			UNCLASSIFIED	1003
9905	79580560 (10957, 10958)			UNCLASSIFIED	1031
9906	10842486 (10963, 10964)			UNCLASSIFIED	1017
9907	77647083 (10965, 10966)			UNCLASSIFIED	1022
9908	86466997 (10975, 10976)			UNCLASSIFIED	1029
9909	33202872 (10977, 10978)			UNCLASSIFIED	1026
9910	46807256 (10989, 10990)			UNCLASSIFIED	1040
9911	65891606 (10993, 10994)			UNCLASSIFIED	1054
9912	30198555 (10995, 10996)			UNCLASSIFIED	1026
9913	5736286 (10997, 10998)			UNCLASSIFIED	1058
9914	32612167 (11001, 11002)			UNCLASSIFIED	1029
9915	65473166 (11003, 11004)			UNCLASSIFIED	1049

9916	54995103 (11005, 11006)			UNCLASSIFIED	1054
9917	30672039 (11013, 11014)			UNCLASSIFIED	1038
9918	17260512 (11037, 11038)			UNCLASSIFIED	1054
9919	17707278 (11039, 11040)			UNCLASSIFIED	1039
9920	35113067 (11051, 11052)			UNCLASSIFIED	1030
9921	56469568 (11057, 11058)			UNCLASSIFIED	1024
9922	17661612 (11061, 11062)			UNCLASSIFIED	1029
9923	25250522 (11067, 11068)			UNCLASSIFIED	1003
9924	80418073 (11073, 11074)			UNCLASSIFIED	1024, 1025
9925	78485858 (11093, 11094)			UNCLASSIFIED	1008
9926	43336027 (11101, 11102)			UNCLASSIFIED	1029
9927	66726728 (11107, 11108)			UNCLASSIFIED	1009
9928	17939048 (11109, 11110)			UNCLASSIFIED	1039
9929	85802525 (11117, 11118)			UNCLASSIFIED	1038
9930	17903125 (11143, 11144)			UNCLASSIFIED	1023
9931	54994379 (11147, 11148)			UNCLASSIFIED	1029
9932	12925101 (11155, 11156)			UNCLASSIFIED	1031
9933	55644235 (11157, 11158)			UNCLASSIFIED	1041

9934	37060371 (11163, 11164)			UNCLASSIFIED	1030
9935	70350240 (11165, 11166)			UNCLASSIFIED	1019
9936	91254980 (11173, 11174)			UNCLASSIFIED	1003
9937	79571321 (11187, 11188)			UNCLASSIFIED	1017
9938	12670831 (11189, 11190)			UNCLASSIFIED	1004, 1022
9939	29452874 (11195, 11196)			UNCLASSIFIED	1024
9940	65465461 (11199, 11200)			UNCLASSIFIED	1038
9941	77883703 (11207, 11208)			UNCLASSIFIED	1021
9942	78793175 (11213, 11214)			UNCLASSIFIED	1008, 1023, 1024
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9944	70824275 (11223, 11224)			UNCLASSIFIED	1011
9945	29144147 (11229, 11230)			UNCLASSIFIED	1017
9946	79468422 (11231, 11232)			UNCLASSIFIED	1009, 1022
9947	27366060 (11237, 11238)			UNCLASSIFIED	1022
9948	79836105 (11247, 11248)			UNCLASSIFIED	1008
9949	8359491 (11251, 11252)			UNCLASSIFIED	1006
9950	66253807 (11253, 11254)			UNCLASSIFIED	1016
9951	9842297 (11255, 11256)			UNCLASSIFIED	1039

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9953	95361599 (11283, 11284)			UNCLASSIFIED	1012
9954	65469305 (11285, 11286)			UNCLASSIFIED	1038
9955	11030810 (11289, 11290)			UNCLASSIFIED	1024
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9957	30530280 (11299, 11300)			UNCLASSIFIED	1026
9958	37056334 (11329, 11330)			UNCLASSIFIED	1030
9959	8756826 (11333, 11334)			UNCLASSIFIED	1004
9960	11617890 (11343, 11344)			UNCLASSIFIED	1004
9961	34873532 (11345, 11346)			UNCLASSIFIED	1029
9962	10030274 (11351, 11352)			UNCLASSIFIED	1003
9963	66365024 (11353, 11354)			UNCLASSIFIED	1004
9964	78471170 (11377, 11378)			UNCLASSIFIED	1044
9965	79865479 (11379, 11380)			UNCLASSIFIED	1017
9966	79425848 (11389, 11390)			UNCLASSIFIED	1017, 1023
9967	33189429 (11425, 11426)			UNCLASSIFIED	1026
9968	30275033 (11433, 11434)			UNCLASSIFIED	1026
9969	10120571 (11439, 11440)			UNCLASSIFIED	1017

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9971	78386473 (11445, 11446)			UNCLASSIFIED	1003
9972	65506547 (11447, 11448)			UNCLASSIFIED	1016
9973	11812368 (11463, 11464)			UNCLASSIFIED	1004
9974	66436580 (11479, 11480)			UNCLASSIFIED	1008, 1024
9975	79574702 (11483, 11484)			UNCLASSIFIED	1031
9976	37060376 (11487, 11488)			UNCLASSIFIED	1030
9977	88098854 (11493, 11494)			UNCLASSIFIED	1025
9978	23298245 (11497, 11498)			UNCLASSIFIED	1044
9979	79782964 (11501, 11502)			UNCLASSIFIED	1017
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9982	20291007 (11525, 11526)			UNCLASSIFIED	1022
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9984	30784153 (11535, 11536)			UNCLASSIFIED	1003
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9987	35017463 (11547, 11548)			UNCLASSIFIED	1054

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9989	65476434 (11561, 11562)			UNCLASSIFIED	1016, 1038
9990	20751678 (11569, 11570)			UNCLASSIFIED	1022
9991	77820032 (11573, 11574)			UNCLASSIFIED	1004, 1024
9992	78681218 (11579, 11580)			UNCLASSIFIED	1003
9993	78786972 (11581, 11582)			UNCLASSIFIED	1008, 1017
9994	6425449 (11593, 11594)			UNCLASSIFIED	1058
9995	78781345 (11599, 11600)			UNCLASSIFIED	1022
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9997	79489587 (11615, 11616)			UNCLASSIFIED	1016
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10005	33700724 (11725, 11726)			UNCLASSIFIED	1034

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10009	10971925 (11749, 11750)			UNCLASSIFIED	1027
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10041	70947156 (12069, 12070)			UNCLASSIFIED	1029

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10044	87896092 (12093, 12094)			UNCLASSIFIED	1001, 1003, 1006, 1008, 1010, 1012, 1015, 1017, 1022, 1023, 1024, 1026, 1027, 1031, 1034, 1038
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10053	78527937 (12175, 12176)			UNCLASSIFIED	1038
10054	81339809 (12179, 12180)			UNCLASSIFIED	1054
10055	11310801 (12185, 12186)			UNCLASSIFIED	1022
10056	79607265 (12187, 12188)			UNCLASSIFIED	1024

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10063	88095683 (12227, 12228)			UNCLASSIFIED	1003, 1008, 1017, 1026, 1034
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10065	55401269 (12241, 12242)			UNCLASSIFIED	1042
10066	11286930 (12245, 12246)			UNCLASSIFIED	1006
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10103	27369871 (12487, 12488)			UNCLASSIFIED	1022
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10105	79175714 (12505, 12506)			UNCLASSIFIED	1003, 1004, 1008, 1016, 1017, 1023, 1024, 1038, 1039, 1044
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10108	78504202 (12515, 12516)			UNCLASSIFIED	1016

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10114	79444624 (12557, 12558)			UNCLASSIFIED	1024
10115	20636759 (12559, 12560)			UNCLASSIFIED	1004
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10177	20295011 (13039, 13040)			UNCLASSIFIED	1022
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10179	14973777 (13047, 13048)			UNCLASSIFIED	1024
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10189	77895258 (13111, 13112)			UNCLASSIFIED	1036
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10192	54700322 (13167, 13168)			UNCLASSIFIED	1022
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10194	66156094 (13173, 13174)			UNCLASSIFIED	1003, 1042
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10197	10169043 (13193, 13194)			UNCLASSIFIED	1003
10198	33102650 (13199, 13200)			UNCLASSIFIED	1026

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10252	21652873 (13621, 13622)			UNCLASSIFIED	1017, 1029, 1058

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10264	17941359 (13721, 13722)			UNCLASSIFIED	1008
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10267	86690753 (13729, 13730)			UNCLASSIFIED	1019
10268	80421407 (13733, 13734)			UNCLASSIFIED	1003, 1010, 1017, 1024, 1025, 1039
10269	25233712 (13737, 13738)			UNCLASSIFIED	1009
10270	46572874 (13751, 13752)			UNCLASSIFIED	1054

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10278	78159374 (13821, 13822)			UNCLASSIFIED	1063
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10281	78404691 (13841, 13842)			UNCLASSIFIED	1016
10282	10340088 (13843, 13844)			UNCLASSIFIED	1038
10283	42563498 (13853, 13854)			UNCLASSIFIED	1000
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10286	79629441 (13867, 13868)			UNCLASSIFIED	1025, 1038
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10288	90934924 (13899, 13900)			UNCLASSIFIED	1020

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10290	95002897 (13915, 13916)			UNCLASSIFIED	1006
10291	20626298 (13937, 13938)			UNCLASSIFIED	1004
10292	79241379 (13943, 13944)			UNCLASSIFIED	1039
10293	65970518 (13953, 13954)			UNCLASSIFIED	1007
10294	65882500 (13955, 13956)			UNCLASSIFIED	1009, 1054
10295	29345835 (13967, 13968)			UNCLASSIFIED	1022
10296	30520815 (13973, 13974)			UNCLASSIFIED	1024
10297	66370541 (13975, 13976)			UNCLASSIFIED	1054
10298	65660033 (13979, 13980)			UNCLASSIFIED	1054
10299	65857534 (13985, 13986)			UNCLASSIFIED	1016
10300	79926139 (13989, 13990)			UNCLASSIFIED	1023
10301	65981117 (13993, 13994)			UNCLASSIFIED	1024
10302	49531052 (14001, 14002)			UNCLASSIFIED	1036
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10305	86685536 (14045, 14046)			UNCLASSIFIED	1029
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10312	51622033 (14137, 14138)			UNCLASSIFIED	1036
10313	77938421 (14139, 14140)			UNCLASSIFIED	1041
10314	100336042 (14145, 14146)			UNCLASSIFIED	1001, 1023, 1053, 1054
10315	34411116 (14151, 14152)			UNCLASSIFIED	1024
10316	54531407 (14157, 14158)			UNCLASSIFIED	1041
10317	27978291 (14163, 14164)			UNCLASSIFIED	1022
10318	27849044 (14171, 14172)			UNCLASSIFIED	1004, 1022, 1034
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10320	80091033 (14181, 14182)			UNCLASSIFIED	1010
10321	23304464 (14187, 14188)			UNCLASSIFIED	1023
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10323	71014802 (14213, 14214)			UNCLASSIFIED	1024
10324	35074443 (14221, 14222)			UNCLASSIFIED	1024

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10328	30420034 (14279, 14280)			UNCLASSIFIED	1016
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10331	37804133 (14309, 14310)			UNCLASSIFIED	1058
10332	78889439 (14319, 14320)			UNCLASSIFIED	1003, 1036
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10335	56069056 (14349, 14350)			UNCLASSIFIED	1021
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10349	28467330 (14461, 14462)			UNCLASSIFIED	1008
10350	38014405 (14473, 14474)			UNCLASSIFIED	1008, 1034
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10353	39549970 (14489, 14490)			UNCLASSIFIED	1038
10354	20622181 (14493, 14494)			UNCLASSIFIED	1003, 1004, 1024, 1034
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10359	79442200 (14515, 14516)			UNCLASSIFIED	1024
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10364	79572942 (14567, 14568)			UNCLASSIFIED	1023, 1044
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10368	8498621 (14593, 14594)			UNCLASSIFIED	1004
10369	78480595 (14617, 14618)			UNCLASSIFIED	1003
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10372	28347999 (14639, 14640)			UNCLASSIFIED	1054
10373	66176875 (14663, 14664)			UNCLASSIFIED	1038
10374	57461624 (14665, 14666)			UNCLASSIFIED	1035
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10377	25131077 (14683, 14684)			UNCLASSIFIED	1008
10378	65875295 (14687, 14688)			UNCLASSIFIED	1044

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10380	79182360 (14699, 14700)			UNCLASSIFIED	1024
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10385	42940996 (14761, 14762)			UNCLASSIFIED	1016, 1050
10386	38057516 (14765, 14766)			UNCLASSIFIED	1054
10387	65660617 (14767, 14768)			UNCLASSIFIED	1053
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10390	80034876 (14803, 14804)			UNCLASSIFIED	1053
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10403	65676603 (14885, 14886)			UNCLASSIFIED	1046
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10405	65878197 (14919, 14920)			UNCLASSIFIED	1038
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10407	79823334 (14977, 14978)			UNCLASSIFIED	1044
10408	66398155 (14979, 14980)			UNCLASSIFIED	1008
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10413	56318078 (15005, 15006)			UNCLASSIFIED	1008
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10418	79566941 (15033, 15034)			UNCLASSIFIED	1003
10419	79621865 (15035, 15036)			UNCLASSIFIED	1025
10420	80078535 (15037, 15038)			UNCLASSIFIED	1022
10421	95012865 (15057, 15058)			UNCLASSIFIED	1001, 1003, 1004, 1006, 1012, 1017, 1024, 1034, 1039, 1054
10422	78976104 (15069, 15070)			UNCLASSIFIED	1030
10423	25139120 (15083, 15084)			UNCLASSIFIED	1026
10424	99079584 (15089, 15090)			UNCLASSIFIED	1004, 1009, 1010, 1016, 1024, 1039, 1049
10425	78486480 (15103, 15104)			UNCLASSIFIED	1009
10426	79554960 (15109, 15110)			UNCLASSIFIED	1022, 1024, 1025
10427	79492377 (15111, 15112)			UNCLASSIFIED	1024
10428	65706518 (15115, 15116)			UNCLASSIFIED	1038
10429	78391719 (15123, 15124)			UNCLASSIFIED	1030
10430	65646524 (15141, 15142)			UNCLASSIFIED	1041

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10433	36514321 (15155, 15156)			UNCLASSIFIED	1029
10434	77541029 (15173, 15174)			UNCLASSIFIED	1053
10435	38322643 (15179, 15180)			UNCLASSIFIED	1033
10436	33700546 (15191, 15192)			UNCLASSIFIED	1026
10437	34115136 (15197, 15198)			UNCLASSIFIED	1000
10438	78250152 (15205, 15206)			UNCLASSIFIED	1029
10439	15024927 (15208, 15210)			UNCLASSIFIED	1054
10440	20442478 (15225, 15226)			UNCLASSIFIED	1024
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10444	65676448 (15251, 15252)			UNCLASSIFIED	1046
10445	30239803 (15253, 15254)			UNCLASSIFIED	1026
10446	11514043 (15261, 15262)			UNCLASSIFIED	1016
10447	87467975 (15269, 15270)			UNCLASSIFIED	1048
10448	65667683 (15281, 15282)			UNCLASSIFIED	1053

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10477	30536017 (15473, 15474)			UNCLASSIFIED	1026
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10480	46892563 (15485, 15486)			UNCLASSIFIED	1034
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10484	34269994 (15503, 15504)			UNCLASSIFIED	1016, 1026

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10489	87896433 (15527, 15528)			UNCLASSIFIED	1015
10490	95290101 (15531, 15532)			UNCLASSIFIED	1004
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10498	10887115 (15605, 15606)			UNCLASSIFIED	1027
10499	78758372 (15609, 15610)			UNCLASSIFIED	1008
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10502	66032451 (15623, 15624)			UNCLASSIFIED	1024, 1053

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10504	86475683 (15653, 15654)			UNCLASSIFIED	1034
10505	57294433 (15657, 15658)			UNCLASSIFIED	1006, 1024, 1034
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10507	27846236 (15663, 15664)			UNCLASSIFIED	1010
10508	87121611 (15665, 15666)			UNCLASSIFIED	1025
10509	10296498 (15667, 15668)			UNCLASSIFIED	1003
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10517	77662342 (15731, 15732)			UNCLASSIFIED	1054
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10520	78461425 (15765, 15766)			UNCLASSIFIED	1026

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10523	79180574 (15807, 15808)			UNCLASSIFIED	1008, 1009, 1022, 1024, 1026, 1044
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10525	78917048 (15815, 15816)			UNCLASSIFIED	1039
10526	8907151 (15831, 15832)			UNCLASSIFIED	1023
10527	65457966 (15833, 15834)			UNCLASSIFIED	1016
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10535	79566837 (15873, 15874)			UNCLASSIFIED	1058
10536	78979309 (15875, 15876)			UNCLASSIFIED	1036, 1054
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10542	78480441 (15905, 15906)			UNCLASSIFIED	1026
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10547	78678187 (15939, 15940)			UNCLASSIFIED	1017
10548	30534224 (15943, 15944)			UNCLASSIFIED	1012
10549	78473961 (15945, 15946)			UNCLASSIFIED	1003
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10551	79852833 (15957, 15958)			UNCLASSIFIED	1017
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10563	79640392 (16055, 16056)			UNCLASSIFIED	1025
10564	27792934 (16075, 16076)			UNCLASSIFIED	1022
10565	16307330 (16083, 16084)			UNCLASSIFIED	1024
10566	79621731 (16095, 16096)			UNCLASSIFIED	1025
10567	35907696 (16113, 16114)			UNCLASSIFIED	1034
10568	11090829 (16115, 16116)			UNCLASSIFIED	1006
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10570	29448234 (16131, 16132)			UNCLASSIFIED	1022
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10574	65462514 (16163, 16164)			UNCLASSIFIED	1023

10575	65982748 (16167, 16168)			UNCLASSIFIED	1004
10576	27976848 (16193, 16194)			UNCLASSIFIED	1022
10577	30409160 (16211, 16212)			UNCLASSIFIED	1026
10578	27844435 (16219, 16220)			UNCLASSIFIED	1022
10579	80249682 (16233, 16234)			UNCLASSIFIED	1003, 1022
10580	79757139 (16235, 16236)			UNCLASSIFIED	1017
10581	79847004 (16241, 16242)			UNCLASSIFIED	1023
10582	21650918 (16245, 16246)			UNCLASSIFIED	1038, 1044
10583	54526968 (16255, 16256)			UNCLASSIFIED	1041
10584	37057654 (16261, 16262)			UNCLASSIFIED	1054
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10588	88096997 (16283, 16284)			UNCLASSIFIED	1025
10589	79596075 (16285, 16286)			UNCLASSIFIED	1003, 1027
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10591	78462606 (16299, 16300)			UNCLASSIFIED	1008
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10595	78095453 (16323, 16324)			UNCLASSIFIED	1021
10596	78260404 (16347, 16348)			UNCLASSIFIED	1029
10597	79762269 (16357, 16358)			UNCLASSIFIED	1008
10598	32699687 (16363, 16364)			UNCLASSIFIED	1000
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10600	43334468 (16385, 16386)			UNCLASSIFIED	1009
10601	21629854 (16393, 16394)			UNCLASSIFIED	1029
10602	20634797 (16403, 16404)			UNCLASSIFIED	1022
10603	17566590 (16405, 16406)			UNCLASSIFIED	1003
10604	79202787 (16425, 16426)			UNCLASSIFIED	1054
10605	65640660 (16437, 16438)			UNCLASSIFIED	1016
10606	65900119 (16439, 16440)			UNCLASSIFIED	1016
10607	79963744 (16447, 16448)			UNCLASSIFIED	1034
10608	78043759 (16465, 16466)			UNCLASSIFIED	1024
10609	17871517 (16475, 16476)			UNCLASSIFIED	1024
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10612	33205261 (16487, 16488)			UNCLASSIFIED	1026
10613	20726424 (16489, 16490)			UNCLASSIFIED	1006, 1034
10614	9882303 (16495, 16496)			UNCLASSIFIED	1008
10615	78793606 (16497, 16498)			UNCLASSIFIED	1026
10616	80500041 (16499, 16500)			UNCLASSIFIED	1003, 1006, 1012, 1024, 1027, 1034
10617	54534616 (16503, 16504)			UNCLASSIFIED	1041
10618	29349432 (16515, 16516)			UNCLASSIFIED	1008
10619	36523786 (16517, 16518)			UNCLASSIFIED	1019
10620	79322040 (16521, 16522)			UNCLASSIFIED	1004
10621	23331987 (16531, 16532)			UNCLASSIFIED	1053
10622	16488222 (16535, 16536)			UNCLASSIFIED	1044
10623	32705516 (16537, 16538)			UNCLASSIFIED	1034
10624	30810733 (16543, 16544)			UNCLASSIFIED	1026
10625	29012146 (16547, 16548)			UNCLASSIFIED	1044
10626	79483144 (16549, 16550)			UNCLASSIFIED	1006
10627	20587078 (16553, 16554)			UNCLASSIFIED	1053
10628	11094215 (16563, 16564)			UNCLASSIFIED	1022

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10631	38068696 (16589, 16590)			UNCLASSIFIED	1009, 1022
10632	50843778 (16599, 16600)			UNCLASSIFIED	1036
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10635	34892205 (16615, 16616)			UNCLASSIFIED	1050
10636	11689021 (16619, 16620)			UNCLASSIFIED	1031
10637	13088374 (16621, 16622)			UNCLASSIFIED	1031
10638	85519672 (16623, 16624)			UNCLASSIFIED	1003, 1004, 1029, 1034, 1044, 1053
10639	79447647 (16635, 16636)			UNCLASSIFIED	1024
10640	65877414 (16639, 16640)			UNCLASSIFIED	1038
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10643	79559077 (16649, 16650)			UNCLASSIFIED	1003
10644	85534541 (16653, 16654)			UNCLASSIFIED	1054
10645	27838048 (16657, 16658)			UNCLASSIFIED	1006
10646	11073159 (16669, 16670)			UNCLASSIFIED	1022

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10649	80025767 (16695, 16696)			UNCLASSIFIED	1006
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10652	30579368 (16711, 16712)			UNCLASSIFIED	1026
10653	10061227 (16717, 16718)			UNCLASSIFIED	1039
10654	66028140 (16719, 16720)			UNCLASSIFIED	1003
10655	29915957 (16737, 16738)			UNCLASSIFIED	1054
10656	35103616 (16761, 16762)			UNCLASSIFIED	1024
10657	9681916 (16763, 16764)			UNCLASSIFIED	1044
10658	78676180 (16765, 16766)			UNCLASSIFIED	1017
10659	79818094 (16775, 16776)			UNCLASSIFIED	1006
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10663	78794010 (16793, 16794)			UNCLASSIFIED	1026
10664	79260934 (16801, 16802)			UNCLASSIFIED	1003

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10666	11308965 (16807, 16808)			UNCLASSIFIED	1024
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10668	78400128 (16837, 16838)			UNCLASSIFIED	1016
10669	78185523 (16847, 16848)			UNCLASSIFIED	1029, 1055
10670	71309207 (16849, 16850)			UNCLASSIFIED	1012, 1029, 1049
10671	86466001 (16859, 16860)			UNCLASSIFIED	1029, 1054
10672	80189508 (16861, 16862)			UNCLASSIFIED	1034, 1038, 1058
10673	29357456 (16867, 16868)			UNCLASSIFIED	1008
10674	65703094 (16893, 16894)			UNCLASSIFIED	1038
10675	46691875 (16913, 16914)			UNCLASSIFIED	1033
10676	79443045 (16949, 16950)			UNCLASSIFIED	1022
10677	66176784 (16951, 16952)			UNCLASSIFIED	1038
10678	77921973 (16967, 16968)			UNCLASSIFIED	1041
10679	10811920 (16979, 16980)			UNCLASSIFIED	1017
10680	54860840 (16985, 16986)			UNCLASSIFIED	1019
10681	77767762 (16991, 16992)			UNCLASSIFIED	1021
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10685	78381247 (17019, 17020)			UNCLASSIFIED	1026
10686	13076318 (17021, 17022)			UNCLASSIFIED	1024
10687	71028028 (17027, 17028)			UNCLASSIFIED	1026
10688	57301098 (17041, 17042)			UNCLASSIFIED	1006
10689	80428023 (17045, 17046)			UNCLASSIFIED	1001, 1003, 1004, 1006, 1008, 1009, 1010, 1012, 1016, 1017, 1022, 1023, 1024, 1025, 1026, 1027, 1031, 1034, 1038, 1039, 1044, 1054
10690	29009947 (17047, 17048)			UNCLASSIFIED	1008
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10692	36512523 (17061, 17062)			UNCLASSIFIED	1026
10693	65488264 (17063, 17064)			UNCLASSIFIED	1023
10694	78392080 (17075, 17076)			UNCLASSIFIED	1030
10695	79418977 (17079, 17080)			UNCLASSIFIED	1034
10696	65465996 (17083, 17084)			UNCLASSIFIED	1022, 1026, 1054

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10698	20617257 (17115, 17116)			UNCLASSIFIED	1004
10699	16405600 (17129, 17130)			UNCLASSIFIED	1003
10700	19510152 (17137, 17138)			UNCLASSIFIED	1034
10701	17939240 (17155, 17156)			UNCLASSIFIED	1038
10702	78798771 (17159, 17160)			UNCLASSIFIED	1009
10703	78675819 (17163, 17164)			UNCLASSIFIED	1022
10704	39562986 (17167, 17168)			UNCLASSIFIED	1006, 1008, 1010, 1017, 1022, 1034
10705	65694586 (17169, 17170)			UNCLASSIFIED	1023
10706	78492224 (17175, 17176)			UNCLASSIFIED	1038
10707	20718851 (17181, 17182)			UNCLASSIFIED	1053
10708	32667859 (17187, 17188)			UNCLASSIFIED	1012, 1022
10709	11392620 (17193, 17194)			UNCLASSIFIED	1024
10710	37003050 (17215, 17216)			UNCLASSIFIED	1001
10711	79260131 (17219, 17220)			UNCLASSIFIED	1024, 1041
10712	66040604 (17221, 17222)			UNCLASSIFIED	1016
10713	23296492 (17223, 17224)			UNCLASSIFIED	1039
10714	78522854 (17225, 17226)			UNCLASSIFIED	1009, 1031

10715	10109197 (17227, 17228)			UNCLASSIFIED	1017
10716	32432208 (17231, 17232)			UNCLASSIFIED	1012
10717	34108349 (17235, 17236)			UNCLASSIFIED	1024
10718	65885952 (17243, 17244)			UNCLASSIFIED	1029
10719	79483804 (17247, 17248)			UNCLASSIFIED	1027
10720	25153403 (17253, 17254)			UNCLASSIFIED	1003
10721	87453870 (17255, 17256)			UNCLASSIFIED	1025
10722	80034523 (17257, 17258)			UNCLASSIFIED	1053
10723	39734888 (17273, 17274)			UNCLASSIFIED	1026, 1029
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10725	57310810 (17277, 17278)			UNCLASSIFIED	1054
10726	46846059 (17295, 17296)			UNCLASSIFIED	1029
10727	49620727 (17297, 17298)			UNCLASSIFIED	1040
10728	39727321 (17303, 17304)			UNCLASSIFIED	1026
10729	13088841 (17305, 17306)			UNCLASSIFIED	1024
10730	65981306 (17313, 17314)			UNCLASSIFIED	1054
10731	21423383 (17345, 17346)			UNCLASSIFIED	1004, 1006, 1022
10732	20746031 (17347, 17348)			UNCLASSIFIED	1010

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10735	66729967 (17379, 17380)			UNCLASSIFIED	1010
10736	79262909 (17397, 17398)			UNCLASSIFIED	1024, 1054
10737	86686329 (17399, 17400)			UNCLASSIFIED	1007
10738	79841192 (17403, 17404)			UNCLASSIFIED	1016, 1017, 1026, 1036
10739	13515365 (17411, 17412)			UNCLASSIFIED	1024
10740	14994467 (17417, 17418)			UNCLASSIFIED	1024
10741	10018834 (17433, 17434)			UNCLASSIFIED	1023
10742	23294762 (17469, 17470)			UNCLASSIFIED	1023
10743	30814342 (17481, 17482)			UNCLASSIFIED	1044
10744	97983937 (17483, 17484)			UNCLASSIFIED	1031
10745	86477681 (17497, 17498)			UNCLASSIFIED	1029
10746	11143689 (17501, 17502)			UNCLASSIFIED	1022
10747	29353096 (17505, 17506)			UNCLASSIFIED	1008
10748	65892209 (17509, 17510)			UNCLASSIFIED	1016
10749	28455563 (17515, 17516)			UNCLASSIFIED	1003, 1008
10750	8490824 (17525, 17526)			UNCLASSIFIED	1034

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10752	32193333 (17529, 17530)			UNCLASSIFIED	1044
10753	39727080 (17533, 17534)			UNCLASSIFIED	1026
10754	39735476 (17543, 17544)			UNCLASSIFIED	1026
10755	17722083 (17549, 17550)			UNCLASSIFIED	1003
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10757	11359343 (17559, 17560)			UNCLASSIFIED	1058
10758	35915893 (17565, 17566)			UNCLASSIFIED	1023
10759	10194473 (17569, 17570)			UNCLASSIFIED	1003
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10762	80046614 (17597, 17598)			UNCLASSIFIED	1024, 1039
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10765	95199342 (17627, 17628)			UNCLASSIFIED	1029
10766	34710708 (17633, 17634)			UNCLASSIFIED	1029
10767	38014671 (17635, 17636)			UNCLASSIFIED	1034
10768	57289067 (17641, 17642)			UNCLASSIFIED	1022

10769	10156717 (17643, 17644)			UNCLASSIFIED	1038
10770	55177172 (17657, 17658)			UNCLASSIFIED	1022
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10772	35992614 (17663, 17664)			UNCLASSIFIED	1034
10773	66582775 (17665, 17666)			UNCLASSIFIED	1009
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10775	20443195 (17681, 17682)			UNCLASSIFIED	1024
10776	35052669 (17683, 17684)			UNCLASSIFIED	1000
10777	94148358 (17685, 17686)			UNCLASSIFIED	1045
10778	79564260 (17703, 17704)			UNCLASSIFIED	1001
10779	66151805 (17713, 17714)			UNCLASSIFIED	1032
10780	9395208 (17749, 17750)			UNCLASSIFIED	1008
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10782	70953360 (17759, 17760)			UNCLASSIFIED	1037
10783	19847725 (17767, 17768)			UNCLASSIFIED	1022
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10785	17727350 (17777, 17778)			UNCLASSIFIED	1044
10786	95287677 (17779, 17780)			UNCLASSIFIED	1012

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10788	71208312 (17789, 17790)			UNCLASSIFIED	1019, 1050
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10790	35067190 (17813, 17814)			UNCLASSIFIED	1054
10791	55478975 (17823, 17824)			UNCLASSIFIED	1026
10792	86477687 (17831, 17832)			UNCLASSIFIED	1029
10793	29361327 (17845, 17846)			UNCLASSIFIED	1023
10794	80232705 (17847, 17848)			UNCLASSIFIED	1022
10795	79824157 (17861, 17862)			UNCLASSIFIED	1017, 1023, 1027, 1038, 1044
10796	57666591 (17867, 17868)			UNCLASSIFIED	1022
10797	17727811 (17873, 17874)			UNCLASSIFIED	1039
10798	87451243 (17895, 17896)			UNCLASSIFIED	1025
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10800	80204559 (17923, 17924)			UNCLASSIFIED	1016, 1023, 1044, 1054
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10802	86469943 (17929, 17930)			UNCLASSIFIED	1029
10803	12921491 (17939, 17940)			UNCLASSIFIED	1004
10804	32554929 (17943, 17944)			UNCLASSIFIED	1006

10805	41086209 (17947, 17948)			UNCLASSIFIED	1037
10806	28305206 (17961, 17962)			UNCLASSIFIED	1054
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10808	37011415 (17965, 17966)			UNCLASSIFIED	1029
10809	94317359 (17971, 17972)			UNCLASSIFIED	1044
10810	11758302 (17985, 17986)			UNCLASSIFIED	1016
10811	10014448 (17997, 17998)			UNCLASSIFIED	1023
10812	11424783 (18001, 18002)			UNCLASSIFIED	1024
10813	78947069 (18015, 18016)			UNCLASSIFIED	1001, 1041, 1054
10814	86665285 (18017, 18018)			UNCLASSIFIED	1019
10815	57052397 (18027, 18028)			UNCLASSIFIED	1023
10816	28977094 (18029, 18030)			UNCLASSIFIED	1026
10817	35893417 (18031, 18032)			UNCLASSIFIED	1053
10818	12829508 (18053, 18054)			UNCLASSIFIED	1031
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10820	17941871 (18071, 18072)			UNCLASSIFIED	1008
10821	70450752 (18077, 18078)			UNCLASSIFIED	1019
10822	78091469 (18091, 18092)			UNCLASSIFIED	1019

10823	65864251 (18099, 18100)			UNCLASSIFIED	1016, 1054
10824	54698345 (18115, 18116)			UNCLASSIFIED	1041
10825	55663467 (18125, 18126)			UNCLASSIFIED	1021
10826	35107119 (18137, 18138)			UNCLASSIFIED	1026
10827	66372542 (18141, 18142)			UNCLASSIFIED	1024
10828	10082755 (18151, 18152)			UNCLASSIFIED	1038
10829	11762730 (18165, 18166)			UNCLASSIFIED	1038
10830	65703434 (18169, 18170)			UNCLASSIFIED	1038
10831	17898192 (18173, 18174)			UNCLASSIFIED	1023
10832	71100385 (18187, 18188)			UNCLASSIFIED	1029
10833	10167011 (18197, 18198)			UNCLASSIFIED	1001
10834	79106566 (18205, 18206)			UNCLASSIFIED	1044
10835	86675527 (18207, 18208)			UNCLASSIFIED	1006, 1007
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10838	78738729 (18217, 18218)			UNCLASSIFIED	1026
10839	94133798 (18221, 18222)			UNCLASSIFIED	1010
10840	79850727 (18233, 18234)			UNCLASSIFIED	1044

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10842	91256447 (18243, 18244)			UNCLASSIFIED	1045
10843	95354590 (18245, 18246)			UNCLASSIFIED	1054
10844	10213248 (18247, 18248)			UNCLASSIFIED	1024
10845	12971964 (18261, 18262)			UNCLASSIFIED	1022, 1024
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10848	65890956 (18291, 18292)			UNCLASSIFIED	1019, 1038
10849	78770238 (18293, 18294)			UNCLASSIFIED	1008
10850	71034263 (18297, 18298)			UNCLASSIFIED	1026
10851	28465641 (18299, 18300)			UNCLASSIFIED	1044
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10853	17931626 (18319, 18320)			UNCLASSIFIED	1023
10854	55652088 (18321, 18322)			UNCLASSIFIED	1012
10855	20403085 (18329, 18330)			UNCLASSIFIED	1004
10856	57530454 (18333, 18334)			UNCLASSIFIED	1004, 1010, 1012, 1022, 1024
10857	65864255 (18361, 18362)			UNCLASSIFIED	1016
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10863	28457354 (18387, 18388)			UNCLASSIFIED	1023
10864	79621081 (18395, 18396)			UNCLASSIFIED	1006, 1009, 1022, 1031
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10866	30171372 (18427, 18428)			UNCLASSIFIED	1030
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10868	66255126 (18431, 18432)			UNCLASSIFIED	1016
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10870	86694351 (18445, 18446)			UNCLASSIFIED	1038
10871	78773042 (18447, 18448)			UNCLASSIFIED	1016, 1054
10872	34406072 (18461, 18462)			UNCLASSIFIED	1054
10873	78077995 (18463, 18464)			UNCLASSIFIED	1029
10874	78079635 (18467, 18468)			UNCLASSIFIED	1054
10875	55778064 (18473, 18474)			UNCLASSIFIED	1042
10876	77653589 (18491, 18492)			UNCLASSIFIED	1053

10877	65458291 (18501, 18502)			UNCLASSIFIED	1024, 1054
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10883	65643368 (18541, 18542)			UNCLASSIFIED	1016
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10885	52496148 (18563, 18564)			UNCLASSIFIED	1036
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10887	46891870 (18571, 18572)			UNCLASSIFIED	1019
10888	9681837 (18575, 18576)			UNCLASSIFIED	1039
10889	80475435 (18585, 18586)			UNCLASSIFIED	1008, 1009, 1012, 1017, 1039
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10891	85805572 (18599, 18600)			UNCLASSIFIED	1038
10892	20292672 (18603, 18604)			UNCLASSIFIED	1034
10893	80085973 (18623, 18624)			UNCLASSIFIED	1054
10894	7417143 (18625, 18626)			UNCLASSIFIED	1058

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10897	78926532 (18655, 18656)			UNCLASSIFIED	1008
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10901	94318981 (18711, 18712)			UNCLASSIFIED	1003
10902	54996140 (18723, 18724)			UNCLASSIFIED	1029
10903	30485699 (18727, 18728)			UNCLASSIFIED	1030
10904	94147136 (18735, 18736)			UNCLASSIFIED	1003
10905	85815026 (18743, 18744)			UNCLASSIFIED	1019
10906	79571889 (18751, 18752)			UNCLASSIFIED	1017
10907	79815513 (18753, 18754)			UNCLASSIFIED	1017
10908	71221510 (18759, 18760)			UNCLASSIFIED	1000
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10911	54534026 (18775, 18776)			UNCLASSIFIED	1041
10912	79777553 (18793, 18794)			UNCLASSIFIED	1017, 1025

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10916	25320134 (18851, 18852)			UNCLASSIFIED	1009
10917	33745823 (18857, 18858)			UNCLASSIFIED	1026
10918	79791976 (18861, 18862)			UNCLASSIFIED	1003, 1017, 1024, 1039
10919	94143130 (18863, 18864)			UNCLASSIFIED	1044
10920	65706751 (18873, 18874)			UNCLASSIFIED	1038
10921	55907795 (18879, 18880)			UNCLASSIFIED	1038
10922	65849072 (18885, 18886)			UNCLASSIFIED	1006
10923	34667089 (18889, 18890)			UNCLASSIFIED	1029
10924	65865577 (18891, 18892)			UNCLASSIFIED	1016
10925	85805747 (18897, 18898)			UNCLASSIFIED	1021, 1054
10926	65486813 (18899, 18900)			UNCLASSIFIED	1054
10927	65893638 (18901, 18902)			UNCLASSIFIED	1038
10928	94240058 (18909, 18910)			UNCLASSIFIED	1003, 1007
10929	79825437 (18919, 18920)			UNCLASSIFIED	1039
10930	79817796 (18925, 18926)			UNCLASSIFIED	1044

10931	28780191 (18939, 18940)			UNCLASSIFIED	1006, 1022
10932	21431991 (18949, 18950)			UNCLASSIFIED	1024
10933	77549185 (18961, 18962)			UNCLASSIFIED	1008
10934	16493398 (18973, 18974)			UNCLASSIFIED	1054
10935	27974931 (18979, 18980)			UNCLASSIFIED	1004
10936	37036825 (18987, 18988)			UNCLASSIFIED	1012
10937	79588075 (18989, 18990)			UNCLASSIFIED	1034
10938	65664168 (18991, 18992)			UNCLASSIFIED	1016
10939	57311306 (18997, 18998)			UNCLASSIFIED	1051
10940	78925408 (19015, 19016)			UNCLASSIFIED	1054
10941	54995480 (19021, 19022)			UNCLASSIFIED	1029
10942	28400540 (19025, 19026)			UNCLASSIFIED	1054
10943	79833819 (19027, 19028)			UNCLASSIFIED	1008
10944	29677376 (19029, 19030)			UNCLASSIFIED	1054
10945	65696399 (19033, 19034)			UNCLASSIFIED	1038, 1054
10946	55504328 (19037, 19038)			UNCLASSIFIED	1021
10947	79454937 (19041, 19042)			UNCLASSIFIED	1016
10948	35893230 (19049, 19050)			UNCLASSIFIED	1053

10949	71048767 (19081, 19082)			UNCLASSIFIED	1037
10950	11690908 (19089, 19090)			UNCLASSIFIED	1016
10951	20438825 (19093, 19094)			UNCLASSIFIED	1010
10952	78935511 (19105, 19106)			UNCLASSIFIED	1017
10953	80218131 (19111, 19112)			UNCLASSIFIED	1003, 1006, 1017, 1022, 1023, 1024, 1026, 1031, 1038, 1039
10954	47660039 (19119, 19120)			UNCLASSIFIED	1029
10955	28344105 (19129, 19130)			UNCLASSIFIED	1024
10956	20477759 (19149, 19150)			UNCLASSIFIED	1034
10957	65481028 (19169, 19170)			UNCLASSIFIED	1049
10958	27981642 (19171, 19172)			UNCLASSIFIED	1006
10959	91235910 (19177, 19178)			UNCLASSIFIED	1010
10960	77713010 (19183, 19184)			UNCLASSIFIED	1054
10961	55890962 (19195, 19196)			UNCLASSIFIED	1017, 1038
10962	11618197 (19203, 19204)			UNCLASSIFIED	1006
10963	79601759 (19207, 19208)			UNCLASSIFIED	1025
10964	24121933 (19213, 19214)			UNCLASSIFIED	1009
10965	65902544 (19223, 19224)			UNCLASSIFIED	1010

10966	38865199 (19233, 19234)			UNCLASSIFIED	1044
10967	66142049 (19239, 19240)			UNCLASSIFIED	1010
10968	52560862 (19245, 19246)			UNCLASSIFIED	1017
10969	95198915 (19251, 19252)			UNCLASSIFIED	1029
10970	30494583 (19255, 19256)			UNCLASSIFIED	1009
10971	25245016 (19257, 19258)			UNCLASSIFIED	1023
10972	78731520 (19287, 19288)			UNCLASSIFIED	1008
10973	29455911 (19291, 19292)			UNCLASSIFIED	1026
10974	38067109 (19293, 19294)			UNCLASSIFIED	1022
10975	20370418 (19299, 19300)			UNCLASSIFIED	1004
10976	79869925 (19301, 19302)			UNCLASSIFIED	1017, 1039
10977	91014540 (19311, 19312)			UNCLASSIFIED	1010
10978	27965794 (19317, 19318)			UNCLASSIFIED	1022
10979	86467870 (19327, 19328)			UNCLASSIFIED	1029
10980	11753830 (19345, 19346)			UNCLASSIFIED	1025
10981	79415370 (19349, 19350)			UNCLASSIFIED	1024
10982	94143136 (19361, 19362)			UNCLASSIFIED	1044
10983	54693589 (19363, 19364)			UNCLASSIFIED	1041

10984	28301771 (19369, 19370)			UNCLASSIFIED	1054
10985	29901155 (19375, 19376)			UNCLASSIFIED	1024
10986	78956188 (19389, 19390)			UNCLASSIFIED	1003
10987	78376065 (19411, 19412)			UNCLASSIFIED	1039
10988	80034539 (19413, 19414)			UNCLASSIFIED	1053
10989	65653165 (19415, 19416)			UNCLASSIFIED	1054
10990	79569807 (19425, 19426)			UNCLASSIFIED	1017
10991	43153913 (19433, 19434)			UNCLASSIFIED	1000
10992	11075820 (19437, 19438)			UNCLASSIFIED	1022
10993	80502762 (19443, 19444)			UNCLASSIFIED	1008, 1012, 1023
10994	20467073 (19449, 19450)			UNCLASSIFIED	1010
10995	65900762 (19451, 19452)			UNCLASSIFIED	1023
10996	65697079 (19457, 19458)			UNCLASSIFIED	1024, 1038
10997	80053616 (19459, 19460)			UNCLASSIFIED	1004
10998	46876354 (19469, 19470)			UNCLASSIFIED	1036
10999	36514357 (19471, 19472)			UNCLASSIFIED	1029
11000	25178491 (19481, 19482)			UNCLASSIFIED	1003
11001	11290575 (19491, 19492)			UNCLASSIFIED	1024

11002	46881479 (19497, 19498)			UNCLASSIFIED	1037
11003	78790087 (19513, 19514)			UNCLASSIFIED	1044
11004	27843641 (19515, 19516)			UNCLASSIFIED	1022
11005	71760804 (19525, 19526)			UNCLASSIFIED	1008, 1009
11006	17941390 (19527, 19528)			UNCLASSIFIED	1008
11007	21414637 (19529, 19530)			UNCLASSIFIED	1006
11008	36825289 (19545, 19546)			UNCLASSIFIED	1054
11009	7423001 (19553, 19554)			UNCLASSIFIED	1058
11010	77877604 (19563, 19564)			UNCLASSIFIED	1019
11011	10814053 (19567, 19568)			UNCLASSIFIED	1003
11012	36902800 (19579, 19580)			UNCLASSIFIED	1026
11013	79750304 (19581, 19582)			UNCLASSIFIED	1010
11014	35107257 (19595, 19596)			UNCLASSIFIED	1030
11015	34104256 (19607, 19608)			UNCLASSIFIED	1024
11016	78467613 (19615, 19616)			UNCLASSIFIED	1008
11017	25326740 (19621, 19622)			UNCLASSIFIED	1003
11018	80047897 (19627, 19628)			UNCLASSIFIED	1022
11019	55310894 (19629, 19630)			UNCLASSIFIED	1042

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11021	17906106 (19845, 19846)			UNCLASSIFIED	1044
11022	39711354 (19863, 19864)			UNCLASSIFIED	1054
11023	30393448 (19677, 19678)			UNCLASSIFIED	1030
11024	78682341 (19683, 19684)			UNCLASSIFIED	1024
11025	37034104 (19685, 19686)			UNCLASSIFIED	1012
11026	38276242 (19691, 19692)			UNCLASSIFIED	1026
11027	46731129 (19693, 19694)			UNCLASSIFIED	1034
11028	65873223 (19699, 19700)			UNCLASSIFIED	1016, 1058
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11030	66255663 (19717, 19718)			UNCLASSIFIED	1016
11031	65661187 (19735, 19736)			UNCLASSIFIED	1041
11032	27848338 (19745, 19746)			UNCLASSIFIED	1010
11033	79183310 (19755, 19756)			UNCLASSIFIED	1024
11034	36902965 (19757, 19758)			UNCLASSIFIED	1026
11035	90831629 (19771, 19772)			UNCLASSIFIED	1000, 1003, 1008, 1022
11036	20399375 (19779, 19780)			UNCLASSIFIED	1004
11037	80073204 (19781, 19782)			UNCLASSIFIED	1001, 1022, 1044

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11040	12809776 (19801, 19802)			UNCLASSIFIED	1031
11041	34128983 (19803, 19804)			UNCLASSIFIED	1026
11042	8018611 (19819, 19820)			UNCLASSIFIED	1022
11043	79556413 (19821, 19822)			UNCLASSIFIED	1023
11044	80218951 (19825, 19826)			UNCLASSIFIED	1003, 1006, 1008, 1024, 1026, 1034
11045	65456240 (19827, 19828)			UNCLASSIFIED	1016, 1038
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11047	79914423 (19835, 19836)			UNCLASSIFIED	1001
11048	94147249 (19843, 19844)			UNCLASSIFIED	1044
11049	9873347 (19845, 19846)			UNCLASSIFIED	1044
11050	55001088 (19847, 19848)			UNCLASSIFIED	1019
11051	19543880 (19851, 19852)			UNCLASSIFIED	1038
11052	94742140 (19861, 19862)			UNCLASSIFIED	1044
11053	79585400 (19873, 19874)			UNCLASSIFIED	1001
11054	55287445 (19875, 19876)			UNCLASSIFIED	1053
11055	29357376 (19891, 19892)			UNCLASSIFIED	1008

11056	35023852 (19893, 19894)			UNCLASSIFIED	1053
11057	39731541 (19895, 19896)			UNCLASSIFIED	1037
11058	10860069 (19897, 19898)			UNCLASSIFIED	1016
11059	42391869 (19907, 19908)			UNCLASSIFIED	1000
11060	16396074 (19911, 19912)			UNCLASSIFIED	1023
11061	46791372 (19951, 19952)			UNCLASSIFIED	1021
11062	25317612 (19953, 19954)			UNCLASSIFIED	1044
11063	46738017 (19965, 19966)			UNCLASSIFIED	1042
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11067	70287213 (19989, 19990)			UNCLASSIFIED	1009
11068	25262598 (20005, 20006)			UNCLASSIFIED	1008
11069	80501497 (20015, 20016)			UNCLASSIFIED	1012
11070	33272462 (20041, 20042)			UNCLASSIFIED	1000
11071	20726725 (20043, 20044)			UNCLASSIFIED	1006
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11073	35101690 (20057, 20058)			UNCLASSIFIED	1024

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11077	79613875 (20087, 20088)			UNCLASSIFIED	1025
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11080	39550238 (20095, 20096)			UNCLASSIFIED	1039
11081	57567486 (20107, 20108)			UNCLASSIFIED	1023
11082	77689694 (20119, 20120)			UNCLASSIFIED	1053
11083	27967525 (20121, 20122)			UNCLASSIFIED	1006
11084	20728342 (20129, 20130)			UNCLASSIFIED	1006
11085	65892227 (20133, 20134)			UNCLASSIFIED	1016
11086	79626465 (20137, 20138)			UNCLASSIFIED	1038
11087	27964527 (20149, 20150)			UNCLASSIFIED	1022
11088	79599522 (20153, 20154)			UNCLASSIFIED	1004
11089	11611543 (20167, 20168)			UNCLASSIFIED	1022, 1024
11090	6380103 (20171, 20172)			UNCLASSIFIED	1058
11091	52205905 (20181, 20182)			UNCLASSIFIED	1036

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11093	86674218 (20195, 20196)			UNCLASSIFIED	1007
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11095	86668708 (20203, 20204)			UNCLASSIFIED	1029, 1034
11096	78235318 (20205, 20206)			UNCLASSIFIED	1026
11097	29020861 (20207, 20208)			UNCLASSIFIED	1008
11098	66435495 (20211, 20212)			UNCLASSIFIED	1034
11099	46861550 (20217, 20218)			UNCLASSIFIED	1050
11100	80237996 (20219, 20220)			UNCLASSIFIED	1034
11101	11024114 (20241, 20242)			UNCLASSIFIED	1004
11102	32059602 (20243, 20244)			UNCLASSIFIED	1026
11103	78527064 (20247, 20248)			UNCLASSIFIED	1008
11104	86464597 (20249, 20250)			UNCLASSIFIED	1029
11105	79779209 (20261, 20262)			UNCLASSIFIED	1017, 1023, 1038
11106	55669564 (20265, 20266)			UNCLASSIFIED	1016
11107	85545152 (20273, 20274)			UNCLASSIFIED	1038
11108	80418100 (20275, 20276)			UNCLASSIFIED	1025
11109	7784522 (20277, 20278)			UNCLASSIFIED	1021

11110	50875041 (20283, 20284)			UNCLASSIFIED	1036
11111	19869659 (20285, 20286)			UNCLASSIFIED	1022
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11114	11084435 (20303, 20304)			UNCLASSIFIED	1024
11115	98177571 (20307, 20308)			UNCLASSIFIED	1012, 1024
11116	78757836 (20315, 20316)			UNCLASSIFIED	1017, 1029
11117	35023442 (20333, 20334)			UNCLASSIFIED	1053
11118	29916038 (20335, 20336)			UNCLASSIFIED	1022
11119	66559129 (20339, 20340)			UNCLASSIFIED	1023, 1024
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11121	58093940 (20347, 20348)			UNCLASSIFIED	1054
11122	5730386 (20349, 20350)			UNCLASSIFIED	1058
11123	13078914 (20351, 20352)			UNCLASSIFIED	1022
11124	33181829 (20359, 20360)			UNCLASSIFIED	1026
11125	78088096 (20365, 20366)			UNCLASSIFIED	1036
11126	54701905 (20367, 20368)			UNCLASSIFIED	1008, 1019
11127	21416447 (20369, 20370)			UNCLASSIFIED	1022

11128	78390961 (20387, 20388)			UNCLASSIFIED	1030
11129	87127037 (20405, 20406)			UNCLASSIFIED	1025
11130	66176078 (20409, 20410)			UNCLASSIFIED	1016
11131	78502459 (20421, 20422)			UNCLASSIFIED	1026
11132	78164313 (20423, 20424)			UNCLASSIFIED	1030
11133	24133062 (20425, 20426)			UNCLASSIFIED	1003
11134	27983392 (20463, 20464)			UNCLASSIFIED	1016
11135	65475114 (20465, 20466)			UNCLASSIFIED	1038
11136	65469088 (20471, 20472)			UNCLASSIFIED	1025, 1054
11137	91230832 (20487, 20488)			UNCLASSIFIED	1003
11138	10840493 (20499, 20500)			UNCLASSIFIED	1017
11139	28979418 (20507, 20508)			UNCLASSIFIED	1039
11140	56512766 (20511, 20512)			UNCLASSIFIED	1019
11141	80504412 (20513, 20514)			UNCLASSIFIED	1001, 1003, 1012, 1017, 1049
11142	54701174 (20525, 20526)			UNCLASSIFIED	1019
11143	78473988 (20529, 20530)			UNCLASSIFIED	1003
11144	78462306 (20549, 20550)			UNCLASSIFIED	1008
11145	78677819 (20573, 20574)			UNCLASSIFIED	1008, 1029

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11148	78496491 (20593, 20594)			UNCLASSIFIED	1038
11149	47660628 (20595, 20596)			UNCLASSIFIED	1050
11150	87454042 (20597, 20598)			UNCLASSIFIED	1025
11151	55185774 (20605, 20606)			UNCLASSIFIED	1019
11152	65329238 (20617, 20618)			UNCLASSIFIED	1057
11153	78745598 (20629, 20630)			UNCLASSIFIED	1028
11154	79864994 (20639, 20640)			UNCLASSIFIED	1006, 1022, 1025, 1038
11155	46739834 (20651, 20652)			UNCLASSIFIED	1034
11156	32444642 (20659, 20660)			UNCLASSIFIED	1024
11157	80230421 (20665, 20666)			UNCLASSIFIED	1003, 1006, 1010, 1016, 1025, 1034, 1054
11158	78087468 (20673, 20674)			UNCLASSIFIED	1029
11159	85806881 (20681, 20682)			UNCLASSIFIED	1054
11160	27778388 (20683, 20684)			UNCLASSIFIED	1004, 1022, 1024
11161	65456754 (20693, 20694)			UNCLASSIFIED	1054
11162	37028812 (20701, 20702)			UNCLASSIFIED	1038
11163	79174016 (20711, 20712)			UNCLASSIFIED	1031

11164	56724149 (20719, 20720)			UNCLASSIFIED	1009, 1031
11165	30257154 (20723, 20724)			UNCLASSIFIED	1024
11166	8722010 (20731, 20732)			UNCLASSIFIED	1053
11167	10263475 (20735, 20736)			UNCLASSIFIED	1023
11168	65886675 (20747, 20748)			UNCLASSIFIED	1016
11169	19638194 (20749, 20750)			UNCLASSIFIED	1054
11170	20430542 (20755, 20756)			UNCLASSIFIED	1004
11171	30268305 (20773, 20774)			UNCLASSIFIED	1008
11172	56518009 (20777, 20778)			UNCLASSIFIED	1008
11173	46571847 (20779, 20780)			UNCLASSIFIED	1029
11174	55455036 (20791, 20792)			UNCLASSIFIED	1016
11175	78958350 (20793, 20794)			UNCLASSIFIED	1016
11176	13083798 (20795, 20796)			UNCLASSIFIED	1027
11177	85817375 (20797, 20798)			UNCLASSIFIED	1019, 1038
11178	16746410 (20799, 20800)			UNCLASSIFIED	1044
11179	15029076 (20803, 20804)			UNCLASSIFIED	1024
11180	78909841 (20813, 20814)			UNCLASSIFIED	1044
11181	42943084 (20819, 20820)			UNCLASSIFIED	1001

11182	86470048 (20825, 20826)			UNCLASSIFIED	1024, 1029
11183	30502164 (20827, 20828)			UNCLASSIFIED	1038
11184	91232298 (20829, 20830)			UNCLASSIFIED	1010
11185	68053558 (20843, 20844)			UNCLASSIFIED	1010, 1026
11186	78381988 (20845, 20846)			UNCLASSIFIED	1023, 1024
11187	65662440 (20859, 20860)			UNCLASSIFIED	1024
11188	78379883 (20865, 20866)			UNCLASSIFIED	1026
11189	80470019 (20869, 20870)			UNCLASSIFIED	1012, 1039
11190	85815208 (20871, 20872)			UNCLASSIFIED	1019
11191	78188559 (20883, 20884)			UNCLASSIFIED	1021, 1022, 1044
11192	16332191 (20887, 20888)			UNCLASSIFIED	1004
11193	65676492 (20889, 20890)			UNCLASSIFIED	1046
11194	16838502 (20897, 20898)			UNCLASSIFIED	1023
11195	30267859 (20903, 20904)			UNCLASSIFIED	1026
11196	78701270 (20907, 20908)			UNCLASSIFIED	1021, 1025
11197	8018293 (20913, 20914)			UNCLASSIFIED	1034
11198	30166581 (20921, 20922)			UNCLASSIFIED	1054
11199	85543256 (20931, 20932)			UNCLASSIFIED	1001, 1038, 1054

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11201	13516619 (20941, 20942)			UNCLASSIFIED	1024
11202	54527129 (20957, 20958)			UNCLASSIFIED	1041
11203	94939037 (20969, 20970)			UNCLASSIFIED	1022, 1034, 1038
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11205	32444158 (20979, 20980)			UNCLASSIFIED	1029
11206	79217362 (20985, 20986)			UNCLASSIFIED	1006
11207	66365404 (20987, 20988)			UNCLASSIFIED	1054
11208	27928083 (20995, 20996)			UNCLASSIFIED	1022
11209	39738002 (20999, 21000)			UNCLASSIFIED	1037
11210	80066308 (21003, 21004)			UNCLASSIFIED	1024
11211	37024263 (21011, 21012)			UNCLASSIFIED	1025
11212	13000752 (21017, 21018)			UNCLASSIFIED	1006, 1024, 1025
11213	65458518 (21027, 21028)			UNCLASSIFIED	1054
11214	80256533 (21029, 21030)			UNCLASSIFIED	1004
11215	35106775 (21065, 21066)			UNCLASSIFIED	1026
11216	27201834 (21073, 21074)			UNCLASSIFIED	1022
11217	17957846 (21077, 21078)			UNCLASSIFIED	1003

11218	77592903 (21079, 21080)			UNCLASSIFIED	1049
11219	10283569 (21101, 21102)			UNCLASSIFIED	1023
11220	56002014 (21105, 21106)			UNCLASSIFIED	1003
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11222	86463651 (21139, 21140)			UNCLASSIFIED	1029
11223	20706034 (21143, 21144)			UNCLASSIFIED	1022
11224	79565159 (21145, 21146)			UNCLASSIFIED	1023
11225	87893277 (21149, 21150)			UNCLASSIFIED	1025
11226	20616850 (21151, 21152)			UNCLASSIFIED	1022
11227	78896959 (21159, 21160)			UNCLASSIFIED	1008
11228	35912877 (21171, 21172)			UNCLASSIFIED	1029
11229	79855069 (21173, 21174)			UNCLASSIFIED	1006, 1017, 1054
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11231	68397206 (21189, 21190)			UNCLASSIFIED	1008
11232	17944820 (21191, 21192)			UNCLASSIFIED	1024
11233	78766036 (21199, 21200)			UNCLASSIFIED	1039
11234	80221555 (21205, 21206)			UNCLASSIFIED	1004, 1022, 1025, 1039
11235	65468331 (21207, 21208)			UNCLASSIFIED	1054

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11237	87116140 (21211, 21212)			UNCLASSIFIED	1018
11238	20608172 (21221, 21222)			UNCLASSIFIED	1022
11239	94847930 (21223, 21224)			UNCLASSIFIED	1044
11240	86473984 (21235, 21236)			UNCLASSIFIED	1054
11241	9897159 (21237, 21238)			UNCLASSIFIED	1023
11242	77674284 (21245, 21246)			UNCLASSIFIED	1054
11243	30420351 (21247, 21248)			UNCLASSIFIED	1044
11244	7541001 (21251, 21252)			UNCLASSIFIED	1058
11245	66027673 (21257, 21258)			UNCLASSIFIED	1003
11246	20287177 (21267, 21268)			UNCLASSIFIED	1022
11247	16835703 (21273, 21274)			UNCLASSIFIED	1023
11248	20467080 (21277, 21278)			UNCLASSIFIED	1010
11249	77882980 (21285, 21286)			UNCLASSIFIED	1042
11250	23294787 (21287, 21288)			UNCLASSIFIED	1003
11251	80577076 (21297, 21298)			UNCLASSIFIED	1034, 1058
11252	85547862 (21301, 21302)			UNCLASSIFIED	1022, 1026, 1053
11253	71092294 (21305, 21306)			UNCLASSIFIED	1053

11254	79114409 (21333, 21334)			UNCLASSIFIED	1016, 1044
11255	56516225 (21369, 21370)			UNCLASSIFIED	1019
11256	10879713 (21375, 21376)			UNCLASSIFIED	1038
11257	38896233 (21379, 21380)			UNCLASSIFIED	1034
11258	65646561 (21385, 21386)			UNCLASSIFIED	1041
11259	34077112 (21389, 21390)			UNCLASSIFIED	1000
11260	8520196 (21391, 21392)			UNCLASSIFIED	1010
11261	65674182 (21393, 21394)			UNCLASSIFIED	1016
11262	65500363 (21395, 21396)			UNCLASSIFIED	1016
11263	79174383 (21397, 21398)			UNCLASSIFIED	1027
11264	55363068 (21399, 21400)			UNCLASSIFIED	1026
11265	65852208 (21401, 21402)			UNCLASSIFIED	1038
11266	77887883 (21429, 21430)			UNCLASSIFIED	1021
11267	65702800 (21433, 21434)			UNCLASSIFIED	1038
11268	70509279 (21437, 21438)			UNCLASSIFIED	1036
11269	78138486 (21439, 21440)			UNCLASSIFIED	1019, 1026
11270	24121945 (21441, 21442)			UNCLASSIFIED	1003
11271	46790037 (21449, 21450)			UNCLASSIFIED	1034

11272	79320737 (21453, 21454)			UNCLASSIFIED	1006
11273	86468597 (21455, 21456)			UNCLASSIFIED	1054
11274	86468696 (21457, 21458)			UNCLASSIFIED	1029
11275	65697190 (21463, 21464)			UNCLASSIFIED	1038
11276	17896584 (21477, 21478)			UNCLASSIFIED	1017
11277	30275096 (21499, 21500)			UNCLASSIFIED	1026
11278	78388849 (21517, 21518)			UNCLASSIFIED	1026
11279	33907355 (21521, 21522)			UNCLASSIFIED	1026
11280	33188289 (21523, 21524)			UNCLASSIFIED	1026
11281	80189534 (21529, 21530)			UNCLASSIFIED	1058
11282	20457001 (21531, 21532)			UNCLASSIFIED	1010
11283	65644292 (21547, 21548)			UNCLASSIFIED	1036, 1041
11284	78778498 (21561, 21562)			UNCLASSIFIED	1001, 1009, 1024, 1054
11285	80069406 (21581, 21582)			UNCLASSIFIED	1010, 1021, 1026
11286	79841684 (21583, 21584)			UNCLASSIFIED	1044
11287	71219765 (21601, 21602)			UNCLASSIFIED	1000, 1024
11288	36830496 (21603, 21604)			UNCLASSIFIED	1054
11289	78924226 (21607, 21608)			UNCLASSIFIED	1008

11290	77944866 (21615, 21616)			UNCLASSIFIED	1029
11291	94321197 (21621, 21622)			UNCLASSIFIED	1003, 1023, 1049
11292	20289207 (21625, 21626)			UNCLASSIFIED	1010
11293	95354098 (21631, 21632)			UNCLASSIFIED	1012, 1016, 1044
11294	78995713 (21633, 21634)			UNCLASSIFIED	1023, 1039, 1044, 1054
11295	34407353 (21641, 21642)			UNCLASSIFIED	1038
11296	78088739 (21643, 21644)			UNCLASSIFIED	1041
11297	52560020 (21645, 21646)			UNCLASSIFIED	1009, 1024
11298	27363055 (21647, 21648)			UNCLASSIFIED	1022
11299	25307387 (21661, 21662)			UNCLASSIFIED	1022
11300	80085445 (21663, 21664)			UNCLASSIFIED	1001, 1008, 1025
11301	55498798 (21669, 21670)			UNCLASSIFIED	1021
11302	65866713 (21675, 21676)			UNCLASSIFIED	1001, 1006, 1044, 1050, 1054
11303	33266314 (21677, 21678)			UNCLASSIFIED	1029
11304	70150053 (21679, 21680)			UNCLASSIFIED	1008, 1016
11305	29144046 (21687, 21688)			UNCLASSIFIED	1026
11306	86685156 (21691, 21692)			UNCLASSIFIED	1019
11307	79830735 (21701, 21702)			UNCLASSIFIED	1023

11308	78180240 (21707, 21708)			UNCLASSIFIED	1024
11309	86476365 (21713, 21714)			UNCLASSIFIED	1029
11310	65697192 (21725, 21726)			UNCLASSIFIED	1038
11311	50243240 (21737, 21738)			UNCLASSIFIED	1022, 1054
11312	78243846 (21741, 21742)			UNCLASSIFIED	1026
11313	33764805 (21743, 21744)			UNCLASSIFIED	1026
11314	19647789 (21745, 21746)			UNCLASSIFIED	1023
11315	80191502 (21749, 21750)			UNCLASSIFIED	1058
11316	8763004 (21751, 21752)			UNCLASSIFIED	1022
11317	94803658 (21765, 21766)			UNCLASSIFIED	1044
11318	28461959 (21767, 21768)			UNCLASSIFIED	1008
11319	94121940 (21769, 21770)			UNCLASSIFIED	1016, 1023
11320	27354378 (21775, 21776)			UNCLASSIFIED	1026
11321	37032500 (21783, 21784)			UNCLASSIFIED	1038
11322	17962386 (21787, 21788)			UNCLASSIFIED	1016
11323	46895889 (21801, 21802)			UNCLASSIFIED	1040
11324	11265667 (21803, 21804)			UNCLASSIFIED	1006
11325	80086651 (21813, 21814)			UNCLASSIFIED	1034

11326	65701668 (21815, 21816)			UNCLASSIFIED	1038
11327	27363573 (21823, 21824)			UNCLASSIFIED	1009, 1022, 1034
11328	10067115 (21837, 21838)			UNCLASSIFIED	1038
11329	65459447 (21841, 21842)			UNCLASSIFIED	1054
11330	90936264 (21857, 21858)			UNCLASSIFIED	1020
11331	34387361 (21861, 21862)			UNCLASSIFIED	1026
11332	57048305 (21865, 21866)			UNCLASSIFIED	1009
11333	54992961 (21867, 21868)			UNCLASSIFIED	1029
11334	79259985 (21879, 21880)			UNCLASSIFIED	1024
11335	47653372 (21901, 21902)			UNCLASSIFIED	1037
11336	77816369 (21905, 21906)			UNCLASSIFIED	1034
11337	20472985 (21907, 21908)			UNCLASSIFIED	1022
11338	77947924 (21909, 21910)			UNCLASSIFIED	1041
11339	54540343 (21915, 21916)			UNCLASSIFIED	1041
11340	33272472 (21917, 21918)			UNCLASSIFIED	1000, 1008
11341	34406471 (21921, 21922)			UNCLASSIFIED	1030, 1038
11342	47652036 (21929, 21930)			UNCLASSIFIED	1029
11343	77811927 (21933, 21934)			UNCLASSIFIED	1054

11344	10206704 (21935, 21936)			UNCLASSIFIED	1025
11345	86466486 (21943, 21944)			UNCLASSIFIED	1029
11346	91228270 (21945, 21946)			UNCLASSIFIED	1010
11347	28382462 (21965, 21966)			UNCLASSIFIED	1044
11348	80071761 (21969, 21970)			UNCLASSIFIED	1022
11349	56333550 (21981, 21982)			UNCLASSIFIED	1019
11350	30402249 (22003, 22004)			UNCLASSIFIED	1008
11351	10114233 (22007, 22008)			UNCLASSIFIED	1038
11352	78206214 (22009, 22010)			UNCLASSIFIED	1022, 1037, 1054
11353	20283462 (22013, 22014)			UNCLASSIFIED	1034
11354	27370022 (22015, 22016)			UNCLASSIFIED	1017
11355	55776948 (22025, 22026)			UNCLASSIFIED	1021
11356	79484394 (22027, 22028)			UNCLASSIFIED	1006
11357	41086158 (22031, 22032)			UNCLASSIFIED	1037
11358	29991100 (22035, 22036)			UNCLASSIFIED	1016
11359	32184184 (22059, 22060)			UNCLASSIFIED	1023
11360	70440943 (22069, 22070)			UNCLASSIFIED	1021, 1041
11361	27963834 (22079, 22080)			UNCLASSIFIED	1010, 1022

11362	79552433 (22093, 22094)			UNCLASSIFIED	1038
11363	79835907 (22095, 22096)			UNCLASSIFIED	1001, 1003, 1008, 1017
11364	17686696 (22115, 22116)			UNCLASSIFIED	1017
11365	25259822 (22127, 22128)			UNCLASSIFIED	1008
11366	37814372 (22129, 22130)			UNCLASSIFIED	1058
11367	20620105 (22137, 22138)			UNCLASSIFIED	1004, 1022
11368	10131332 (22141, 22142)			UNCLASSIFIED	1039
11369	56139990 (22151, 22152)			UNCLASSIFIED	1009
11370	91230842 (22181, 22182)			UNCLASSIFIED	1010
11371	20553107 (22189, 22190)			UNCLASSIFIED	1053
11372	78762686 (22191, 22192)			UNCLASSIFIED	1030
11373	23282436 (22193, 22194)			UNCLASSIFIED	1044
11374	46887740 (22197, 22198)			UNCLASSIFIED	1037
11375	20297488 (22199, 22200)			UNCLASSIFIED	1034
11376	79208839 (22207, 22208)			UNCLASSIFIED	1001
11377	30258872 (22209, 22210)			UNCLASSIFIED	1008
11378	10066879 (22211, 22212)			UNCLASSIFIED	1025
11379	78700415 (22219, 22220)			UNCLASSIFIED	1008, 1009

11380	65507923 (22227, 22228)			UNCLASSIFIED	1038
11381	81637048 (22229, 22230)			UNCLASSIFIED	1054
11382	80059002 (22237, 22238)			UNCLASSIFIED	1022, 1036
11383	56513371 (22249, 22250)			UNCLASSIFIED	1019
11384	20618789 (22253, 22254)			UNCLASSIFIED	1004
11385	65661997 (22267, 22268)			UNCLASSIFIED	1010, 1016
11386	20723097 (22283, 22284)			UNCLASSIFIED	1024
11387	98071577 (22291, 22292)			UNCLASSIFIED	1034
11388	23307117 (22293, 22294)			UNCLASSIFIED	1053
11389	79427790 (22295, 22296)			UNCLASSIFIED	1017
11390	17681027 (22297, 22298)			UNCLASSIFIED	1017
11391	27366670 (22299, 22300)			UNCLASSIFIED	1022
11392	46694910 (22307, 22308)			UNCLASSIFIED	1034
11393	34384284 (22309, 22310)			UNCLASSIFIED	1054
11394	85805936 (22315, 22316)			UNCLASSIFIED	1054
11395	71092305 (22319, 22320)			UNCLASSIFIED	1053
11396	87453071 (22321, 22322)			UNCLASSIFIED	1025
11397	78376391 (22331, 22332)			UNCLASSIFIED	1026

11398	79186451 (22337, 22338)			UNCLASSIFIED	1027
11399	7431752 (22346, 22346)			UNCLASSIFIED	1058
11400	10173601 (22351, 22352)			UNCLASSIFIED	1023
11401	28965795 (22353, 22354)			UNCLASSIFIED	1044
11402	65884225 (22355, 22356)			UNCLASSIFIED	1054
11403	21425260 (22359, 22360)			UNCLASSIFIED	1022
11404	11245692 (22369, 22370)			UNCLASSIFIED	1022, 1034
11405	34899667 (22371, 22372)			UNCLASSIFIED	1054
11406	33765171 (22373, 22374)			UNCLASSIFIED	1026
11407	57308564 (22377, 22378)			UNCLASSIFIED	1010
11408	39444106 (22383, 22384)			UNCLASSIFIED	1022
11409	35938636 (22385, 22386)			UNCLASSIFIED	1023
11410	10866089 (22391, 22392)			UNCLASSIFIED	1016
11411	65682366 (22395, 22396)			UNCLASSIFIED	1041
11412	10153872 (22411, 22412)			UNCLASSIFIED	1025
11413	30256320 (22449, 22450)			UNCLASSIFIED	1024
11414	65706652 (22451, 22452)			UNCLASSIFIED	1038
11415	82116243 (22453, 22454)			UNCLASSIFIED	1016, 1017

11416	33183160 (22459, 22460)			UNCLASSIFIED	1026
11417	51577850 (22461, 22462)			UNCLASSIFIED	1036
11418	20485265 (22463, 22464)			UNCLASSIFIED	1022
11419	21416819 (22471, 22472)			UNCLASSIFIED	1034
11420	79563885 (22473, 22474)			UNCLASSIFIED	1001
11421	37799373 (22475, 22476)			UNCLASSIFIED	1012
11422	35900314 (22481, 22482)			UNCLASSIFIED	1053
11423	52468781 (22493, 22494)			UNCLASSIFIED	1036
11424	29142422 (22509, 22510)			UNCLASSIFIED	1017
11425	25257349 (22515, 22516)			UNCLASSIFIED	1026
11426	78721360 (22525, 22526)			UNCLASSIFIED	1026
11427	80190347 (22529, 22530)			UNCLASSIFIED	1009, 1058
11428	77868173 (22531, 22532)			UNCLASSIFIED	1041
11429	77657253 (22541, 22542)			UNCLASSIFIED	1053
11430	94139658 (22551, 22552)			UNCLASSIFIED	1022, 1044
11431	86684236 (22563, 22564)			UNCLASSIFIED	1029
11432	21395374 (22565, 22566)			UNCLASSIFIED	1022
11433	29012863 (22573, 22574)			UNCLASSIFIED	1017

11434	65471939 (22577, 22578)			UNCLASSIFIED	1054
11435	79207587 (22579, 22580)			UNCLASSIFIED	1026, 1054
11436	78749475 (22581, 22582)			UNCLASSIFIED	1008
11437	65987177 (22583, 22584)			UNCLASSIFIED	1054
11438	87115329 (22595, 22596)			UNCLASSIFIED	1025
11439	8489553 (22597, 22598)			UNCLASSIFIED	1024
11440	66123144 (22601, 22602)			UNCLASSIFIED	1003
11441	52561506 (22619, 22620)			UNCLASSIFIED	1003
11442	8521991 (22627, 22628)			UNCLASSIFIED	1034
11443	78764847 (22635, 22636)			UNCLASSIFIED	1022, 1026, 1044
11444	30788047 (22639, 22640)			UNCLASSIFIED	1026
11445	65883040 (22645, 22646)			UNCLASSIFIED	1054
11446	17888579 (22655, 22656)			UNCLASSIFIED	1023
11447	28810559 (22657, 22658)			UNCLASSIFIED	1044
11448	95196868 (22663, 22664)			UNCLASSIFIED	1054
11449	37015827 (22665, 22666)			UNCLASSIFIED	1029
11450	54532171 (22679, 22680)			UNCLASSIFIED	1041
11451	86380123 (22683, 22684)			UNCLASSIFIED	1053

11452	66465961 (22691, 22692)			UNCLASSIFIED	1024
11453	94830591 (22715, 22716)			UNCLASSIFIED	1034, 1038
11454	78491961 (22725, 22726)			UNCLASSIFIED	1054
11455	85540325 (22733, 22734)			UNCLASSIFIED	1038
11456	79262949 (22735, 22736)			UNCLASSIFIED	1039
11457	16410791 (22737, 22738)			UNCLASSIFIED	1030
11458	23283217 (22739, 22740)			UNCLASSIFIED	1039
11459	80079428 (22747, 22748)			UNCLASSIFIED	1034
11460	78791981 (22755, 22756)			UNCLASSIFIED	1009
11461	20621522 (22761, 22762)			UNCLASSIFIED	1006
11462	20623368 (22767, 22768)			UNCLASSIFIED	1022
11463	30639142 (22779, 22780)			UNCLASSIFIED	1022
11464	27966827 (22785, 22786)			UNCLASSIFIED	1053
11465	56518019 (22793, 22794)			UNCLASSIFIED	1008
11466	87458665 (22799, 22800)			UNCLASSIFIED	1048
11467	55183764 (22807, 22808)			UNCLASSIFIED	1019
11468	15019433 (22813, 22814)			UNCLASSIFIED	1054
11469	19506747 (22819, 22820)			UNCLASSIFIED	1038

11470	27836166 (22821, 22822)			UNCLASSIFIED	1006
11471	77863285 (22823, 22824)			UNCLASSIFIED	1021
11472	85800972 (22833, 22834)			UNCLASSIFIED	1025, 1053
11473	77551208 (22841, 22842)			UNCLASSIFIED	1054
11474	54528438 (22849, 22850)			UNCLASSIFIED	1041
11475	35105815 (22857, 22858)			UNCLASSIFIED	1029
11476	66405218 (22863, 22864)			UNCLASSIFIED	1024
11477	80483946 (22865, 22866)			UNCLASSIFIED	1012
11478	85520663 (22869, 22870)			UNCLASSIFIED	1049
11479	86467265 (22879, 22880)			UNCLASSIFIED	1029
11480	80484082 (22885, 22886)			UNCLASSIFIED	1012, 1029, 1034
11481	66152855 (22889, 22890)			UNCLASSIFIED	1003
11482	77674101 (22891, 22892)			UNCLASSIFIED	1054
11483	27846452 (22895, 22896)			UNCLASSIFIED	1022
11484	29018420 (22897, 22898)			UNCLASSIFIED	1044
11485	65890995 (22907, 22908)			UNCLASSIFIED	1054
11486	19646090 (22917, 22918)			UNCLASSIFIED	1004
11487	80105359 (22919, 22920)			UNCLASSIFIED	1053

11488	79574895 (22935, 22936)			UNCLASSIFIED	1031
11489	78424663 (22943, 22944)			UNCLASSIFIED	1030
11490	78037099 (22945, 22946)			UNCLASSIFIED	1050
11491	57055428 (22963, 22964)			UNCLASSIFIED	1051

TABLE 2.

TISSUE ID NUMBERS FROM 1000 - 1058

TRA 1521931v1

tissue id	tissue heirarchy	Common conditions/diseases	Other diseases
1000	Cardiovascular System/Artery/Coronary Artery	cancer,trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Atherosclerosis, Aneurysm, Hypertension, Fibromuscular dysplasia, Stroke, Scleroderma,Obesity,Transplantation, Myocardial infarction, Embolism, Cardiovascular disorders, Bypass surgery,
1001	Cardiovascular System/Heart	cancer,trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Cardiomyopathy, Atherosclerosis,Hypertension, Congenital heart defects, Aortic stenosis ,Atrial septal defect (ASD),Atrioventricular (A-V) canal defect, Ductus arteriosus , Pulmonary stenosis , Subaortic stenosis, Ventricular septal defect (VSD), valve diseases,Tuberous sclerosis, Scleroderma, Obesity, Transplantation
1002	Cardiovascular System/Vein/Umbilical Vein	cancer,trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Anemia , Bleeding disorders, Scleroderma, Transplantation
1003	Endocrine System/Adrenal Gland/Suprarenal gland	cancer,trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Adrenoleukodystrophy , Congenital Adrenal Hyperplasia,
1004	Endocrine System/Pancreas	cancer,trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Diabetes,Von Hippel-Lindau (VHL) syndrome , Pancreatitis,Obesity
1005	Endocrine System/Parathyroid Gland	cancer,trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Hyperparathyroidism, Hypoparathyroidism
1006	Endocrine System/Thyroid	cancer,trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Hypothyroidism and Hypothyroidism
1007	Female Reproductive System/Cervix	cancer,trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Fertility
1008	Female Reproductive System/Mammary gland/Breast	cancer,trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Fertility
1009	Female Reproductive System/Ovary	cancer,trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Endometriosis,Fertility
1010	Female Reproductive System/Placenta	cancer,trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Fertility
1011	Female Reproductive System/Placenta/Chorionic Villus	cancer,trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Fertility
1012	Female Reproductive System/Uterus	cancer,trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Endometriosis,Fertility

1013	Female Reproductive System/Uterus/Myometrium	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Endometriosis, Fertility
1014	Gastro-intestinal/Digestive System/Intestine/Colon	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Hirschsprung's disease, Crohn's Disease, Appendicitis
1015	Gastro-intestinal/Digestive System/Large Intestine/Colon/Ascending Colon	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Hirschsprung's disease, Crohn's Disease, Appendicitis
1016	Gastro-intestinal/Digestive System/Liver	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Von Hippel-Lindau (VHL) syndrome, Cirrhosis, Transplantation
1017	Gastro-intestinal/Digestive System/Salivary Glands	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Xerostomia
1018	Gastro-intestinal/Digestive System/Salivary Glands/Parotid Salivary glands	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Xerostomia
1019	Gastro-intestinal/Digestive System/Small Intestine	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Inflammatory bowel disease, Diverticular disease,
1020	Gastro-intestinal/Digestive System/Small Intestine/Duodenum	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Inflammatory bowel disease, Diverticular disease,
1021	Gastro-intestinal/Digestive System/Stomach	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Hypercalcemia, Ulcers
1022	Hematopoietic and Lymphatic System/Hematopoietic Tissues/Bone Marrow	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Hemophilia, hypercoagulation, idiopathic thrombocytopenic purpura, autoimmune disease, allergies, immunodeficiencies, transplantation, Graft versus host,
1023	Hematopoietic and Lymphatic System/Hematopoietic Tissues/Lymphoid tissue	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Hemophilia, hypercoagulation, idiopathic thrombocytopenic purpura, autoimmune disease, allergies, immunodeficiencies, transplantation, Graft versus host disease (GVHD), Lymphoedema
1024	Hematopoietic and Lymphatic System/Hematopoietic Tissues/Lymphoid tissue/Lymph node	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Lymphoedema, Allergies
1025	Hematopoietic and Lymphatic System/Hematopoietic Tissues/Lymphoid tissue/Spleen	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Hemophilia, Hypercoagulation, Idiopathic thrombocytopenic purpura, immunodeficiencies, Graft versus host

1026	Hematopoietic and Lymphatic System/Hematopoietic Tissues/Peripheral Blood	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Anemia, Ataxia-telangiectasia, Autoimmune disease, Immunodeficiencies
1027	Hematopoietic and Lymphatic System/Hematopoietic Tissues/Thymus	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Hemophilia, hypercoagulation, Idiopathic thrombocytopenic purpura, Immunodeficiencies
1028	Hematopoietic and Lymphatic System/Hematopoietic Tissues/Tonsils	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Tonsillitis
1029	Male Reproductive System/Prostate	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Fertility
1030	Male Reproductive System/Testis	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Fertility
1031	Musculoskeletal System/Bone	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Osteoporosis, Hypercalcemia, Arthritis, Ankylosing spondylitis, Scoliosis
1032	Musculoskeletal System/Muscle	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Muscular dystrophy, Lesch-Nyhan syndrome, Myasthenia gravis
1033	Musculoskeletal System/Muscle/Skeletal Muscle	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Muscular dystrophy, Lesch-Nyhan syndrome, Myasthenia gravis
1034	Nervous System/Brain	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Von Hippel-Lindau (VHL) syndrome, Alzheimer's disease, Stroke, Tuberous sclerosis, hypercalcemia, Parkinson's disease, Huntington's disease, Cerebral palsy, Epilepsy, Lesch-Nyhan syndrome, Multiple sclerosis, Ataxia-telangiectasia, Leukodystrophies, Behavioral disorders, Addiction, Anxiety, Pain, Neuroprotection
1035	Nervous System/Brain/Brain Stem/Hindbrain/Cerebellum	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Von Hippel-Lindau (VHL) syndrome, Alzheimer's disease, Stroke, Tuberous sclerosis, hypercalcemia, Parkinson's disease, Huntington's disease, Cerebral palsy, Epilepsy, Lesch-Nyhan syndrome, Multiple sclerosis, Ataxia-telangiectasia, Leukodystrophies, Behavioral disorders, Addiction, Anxiety, Pain, Neuroprotection

1036	Nervous System/Brain/Hippocampus	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Von Hippel-Lindau (VHL) syndrome, Alzheimer's disease, Stroke, Tuberous sclerosis, hypercalcaemia, Parkinson's disease, Huntington's disease, Cerebral palsy, Epilepsy, Lesch-Nyhan syndrome, Multiple sclerosis, Ataxia-telangiectasia, Leukodystrophies, Behavioral disorders, Addiction, Anxiety, Pain, Neuroprotection
1037	Nervous System/Brain/Prosencephalon/Forebrain/Diencephalon/Hypothalamus	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Endocrine dysfunctions, Diabetes, obesity, Growth and reproductive disorders
1038	Nervous System/Brain/Prosencephalon/Forebrain/Diencephalon/Pituitary Gland	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Endocrine dysfunctions, Diabetes, obesity, Growth and reproductive disorders
1039	Nervous System/Brain/Prosencephalon/Forebrain/Diencephalon/Thalamus	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Von Hippel-Lindau (VHL) syndrome, Alzheimer's disease, Stroke, Tuberous sclerosis, hypercalcaemia, Parkinson's disease, Huntington's disease, Cerebral palsy, Epilepsy, Lesch-Nyhan syndrome, Multiple sclerosis, Ataxia-telangiectasia, Leukodystrophies, Behavioral disorders, Addiction, Anxiety, Pain, Neuroprotection
1040	Nervous System/Brain/Prosencephalon/Forebrain/Telencephalon/Cerebrum/Cerebral Cortex/Cerebral Gray Matter/Cerebral gyri/Temporal Lobe/Amygdala	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Von Hippel-Lindau (VHL) syndrome, Alzheimer's disease, Stroke, Tuberous sclerosis, hypercalcaemia, Parkinson's disease, Huntington's disease, Cerebral palsy, Epilepsy, Lesch-Nyhan syndrome, Multiple sclerosis, Ataxia-telangiectasia, Leukodystrophies, Behavioral disorders, Addiction, Anxiety, Pain, Neuroprotection

1041	Nervous System/Brain/Substantia Nigra	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Von Hippel-Lindau (VHL) syndrome , Alzheimer's disease, Stroke, Tuberous sclerosis, hypercalcemia, Parkinson's disease, Huntington's disease, Cerebral palsy, Epilepsy, Lesch-Nyhan syndrome, Multiple sclerosis, Ataxia-telangiectasia, Leukodystrophies, Behavioral disorders, Addiction, Anxiety, Pain, Neuroprotection
1042	Nervous System/Spinal Chord	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Multiple sclerosis, Leukodystrophies, Pain, Neuroprotection
1043	Respiratory System/Larynx	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Laryngitis, Asthma
1044	Respiratory System/Lung	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Systemic lupus erythematosus , Autoimmune disease, Asihma, Emphysema, Scleroderma
1045	Respiratory System/Lung/Lung Pleura	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Systemic lupus erythematosus , Autoimmune disease, Asihma, Emphysema, Scleroderma, allergy, ARDS.
1046	Respiratory System/Trachea	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Asthma, allergy, ARDS,
1047	Sensory System/Ear/Bony Labyrinth of inner ear/Cochlea	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Hearing loss, Tinnitus
1048	Sensory System/Skin	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Psoriasis, Actinic keratosis , Tuberous sclerosis, Acne, Hair growth, alopecia, pigmentation disorders, endocrine disorders
1049	Sensory System/Skin/Dermis	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Psoriasis, Actinic keratosis , Tuberous sclerosis, Acne, Hair growth, alopecia, pigmentation disorders, endocrine disorders
1050	Sensory System/Skin/Epidermis	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Psoriasis, Actinic keratosis , Tuberous sclerosis, Acne, Hair growth, alopecia, pigmentation disorders, endocrine disorders
1051	Sensory System/Skin/Epidermis/Hair Follicles	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Psoriasis, Actinic keratosis , Tuberous sclerosis, Acne, Hair growth, alopecia, pigmentation disorders, endocrine disorders
1052	Sensory System/Skin/Foreskin	cancer, trauma, regeneration (in vitro and in vivo), viral/bacterial/parasitic infections	Psoriasis, Actinic keratosis , Tuberous sclerosis
1053	Tissue Not Assigned		

What is claimed is:

1. An isolated nucleic acid molecule encoding a polypeptide comprising an amino acid sequence that is at least 85% identical to a polypeptide including an amino acid sequence selected from the group consisting of SEQ ID NO: 4988, 4990, 4992, 4994, 4996, 4998, 5000, 5076, 5078, 5080, 19474, 19476, 19478, 11740, 11742 and 11744, or the complement thereof.
2. The isolated nucleic acid molecule of claim 1, said molecule hybridizing under stringent conditions to a nucleic acid sequence complementary to a nucleic acid molecule comprising the sequence of nucleotides selected from the group consisting of SEQ ID NO. 4987, 4989, 4991, 4993, 4995, 4997, 4999, 5075, 5077, 5079, 19473, 19475, 19477, 11739, 11741 and 11743, or the complement thereof.
3. The isolated nucleic acid molecule of claim 1, said molecule encoding a polypeptide comprising the amino acid sequence selected from the group consisting of SEQ ID NO: 4988, 4990, 4992, 4994, 4996, 4998, 5000, 5076, 5078, 5080, 19474, 19476, 19478, 11740, 11742 and 11744, or an amino acid sequence comprising one or more conservative substitutions in the amino acid sequence selected from the group consisting of SEQ ID NO: 4988, 4990, 4992, 4994, 4996, 4998, 5000, 5076, 5078, 5080, 19474, 19476, 19478, 11740, 11742 and 11744.
4. The isolated nucleic acid molecule of claim 1, wherein said molecule encodes a polypeptide comprising the amino acid sequence selected from the group consisting of SEQ ID NO: 4988, 4990, 4992, 4994, 4996, 4998, 5000, 5076, 5078, 5080, 19474, 19476, 19478, 11740, 11742 and 11744.
5. The isolated nucleic acid molecule of claim 1, wherein said molecule comprises the sequence of nucleotides selected from the group consisting of SEQ ID NO. 4987, 4989, 4991, 4993, 4995, 4997, 4999, 5075, 5077, 5079, 19473, 19475, 19477, 11739, 11741 and 11743, or the complement thereof.

6. An oligonucleotide less than 100 nucleotides in length and comprising at least 6 contiguous nucleotides selected from the group consisting of SEQ ID NO. 4987, 4989, 4991, 4993, 4995, 4997, 4999, 5075, 5077, 5079, 19473, 19475, 19477, 11739, 11741 and 11743, or the complement thereof.
7. A vector comprising the nucleic acid molecule of claim 1.
8. The vector of claim 7, wherein said vector is an expression vector.
9. A host cell comprising the isolated nucleic acid molecule of claim 1.
10. A substantially purified polypeptide comprising an amino acid sequence at least 80% identical to a polypeptide comprising the amino acid sequence selected from the group consisting of SEQ ID NO: 4988, 4990, 4992, 4994, 4996, 4998, 5000, 5076, 5078, 5080, 19474, 19476, 19478, 11740, 11742 and 11744.
11. The polypeptide of claim 10, wherein said polypeptide comprises the amino acid sequence selected from the group consisting of SEQ ID NO: 4988, 4990, 4992, 4994, 4996, 4998, 5000, 5076, 5078, 5080, 19474, 19476, 19478, 11740, 11742 and 11744.
12. An antibody that selectively binds to the polypeptide of claim 10.
13. A pharmaceutical composition comprising a therapeutically or prophylactically effective amount of a therapeutic selected from the group consisting of:
 - a) the nucleic acid of claim 1;
 - b) the polypeptide of claim 10; and
 - c) the antibody of claim 12;and a pharmaceutically acceptable carrier.

14. A kit comprising in one or more containers, a therapeutically or prophylactically effective amount of the pharmaceutical composition of claim 13.

15. A method of producing the polypeptide of claim 10, said method comprising culturing the host cell of claim 9 under conditions in which the nucleic acid molecule is expressed.

16. A method of detecting the presence of the polypeptide of claim 10 in a sample, comprising contacting the sample with a compound that selectively binds to said polypeptide under conditions allowing the formation of a complex between said polypeptide and said compound, and detecting said complex, if present, thereby identifying said polypeptide in said sample.

17. A method of detecting the presence of a nucleic acid molecule of claim 1 in a sample, the method comprising contacting the sample with a nucleic acid probe or primer that selectively binds to the nucleic acid molecule and determining whether the nucleic acid probe or primer bound to the nucleic acid molecule of claim 1 is present in the sample.

18. A method for modulating the activity of the polypeptide of claim 10, the method comprising contacting a cell sample comprising the polypeptide of claim 10 with a compound that binds to said polypeptide in an amount sufficient to modulate the activity of the polypeptide.

19. The use of a therapeutic in the manufacture of a medicament for treating a syndrome associated with a ORFX-associated disorder, wherein said therapeutic is selected from the group consisting of:

- a) the nucleic acid of claim 1;
- b) the polypeptide of claim 10; and
- c) the antibody of claim 12.

20. A method for screening for a modulator of activity or of latency or predisposition to an ORFX-associated disorder, said method comprising:

- a) contacting a test compound with the polypeptide of claim 10; and
- b) determining if said test compound binds to said polypeptide,

wherein binding of said test compound to said polypeptide indicates the test compound is a modulator of activity or of latency or predisposition to an ORFX-associated disorder.

21. A method for screening for a modulator of activity or of latency or predisposition to an ORFX-associated disorder, said method comprising:

- a) administering a test compound to a test subject at an increased risk ORFX-associated disorder, wherein said test subject recombinantly expresses a polypeptide encoded by the nucleotide of claim 1;
- b) measuring expression the activity of said protein in said test subject;
- c) measuring the activity of said protein in a control subject that recombinantly expresses said protein and is not at increased risk for an ORFX-associated disorder; and
- d) comparing expression of said protein in said test subject and said control subject, wherein a change in the activity of said protein in said test subject relative to said control subject indicates the test compound is a modulator or of latency of predisposition to an ORFX-associated disorder.

22. The method of claim 20, wherein said test animal is a recombinant test animal that expresses a test protein transgene or expresses said transgene under the control of a promoter at an increased level relative to a wild-type test animal, and wherein said promoter is not the native gene promoter of said transgene.

23. A method for determining the presence of or predisposition to a disease associated with altered levels of a polypeptide of claim 11 in a subject, the method comprising:

- a) measuring the amount of the polypeptide in a sample from said subject; and

b) comparing the amount of said polypeptide in step (a) to the amount of the polypeptide present in a control sample,
wherein an alteration in the level of the polypeptide in step (a) as compared to the control sample indicates the presence of or predisposition to a disease in said subject.

24. The method of claim 23, wherein said subject is a human.

25. A method for determining the presence of or predisposition to a disease associated with altered levels the nucleic acid molecule of claim 1 in a subject, the method comprising:

- a) measuring the amount of the nucleic acid in a sample from the mammalian subject;
and
- b) comparing the amount of said nucleic acid in step (a) to the amount of the nucleic acid present in a control sample,

wherein an alteration in the level of the nucleic acid in step (a) as compared to the control sample indicates the presence of or predisposition to said disease in said subject.

26. The method of claim 25, wherein said subject is a human.

27. A method of treating or preventing a pathological condition associated with an ORFX-associated disorder in a subject, the method comprising administering to said subject the polypeptide of claim 10 in an amount sufficient to alleviate or prevent said pathological condition.

28. The method of claim 27, wherein said subject is a human.

29. A method of treating or preventing a pathological condition associated with an ORFX-associated disorder in a subject, the method comprising administering to said subject the nucleic acid molecule of claim 1 in an amount sufficient to alleviate or prevent said pathological condition.

30. The method of claim 29, wherein said subject is a human.

31. A method of treating or preventing a pathological condition associated with an ORFX-associated disorder in a subject, the method comprising administering to said subject the antibody of claim 12 in an amount sufficient to alleviate or prevent said pathological condition.

32. The method of claim 31, wherein said subject is a human.